GenCore version 5.1.6

OM nucleic - nucleic search, using sw model

Run on: August 17, 2005, 17:23:27 ; Search time 2446.54 Seconds
(without alignments)
11060.388 Million cell updates/sec

Title: US-09-996-630A-11
Perfect score: 4165
Sequence: 1 gtctaaagagtgtaaagacc.....acatattggtgttcagcaat 4165

Scoring table: IDENTITY_NUC
Gapop 10.0 , Gapext 1.0

Searched: 7316285 seqs, 3248459403 residues

Total number of hits satisfying chosen parameters: 14632570

Minimum DB seq length: 0
Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries

Database : Published_Applications_NA:*
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

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1	4165	100.0	4165	10	US-09-873-367C-95	Sequence 95, Appl
2	4165	100.0	4165	21	US-10-843-641A-95	Sequence 95, Appl
3	4163.4	100.0	5519	14	US-10-097-340-136	Sequence 136, App
4	4163.4	100.0	5519	16	US-10-354-358-69	Sequence 69, Appl
5	4138.2	99.4	4176	18	US-10-425-114-26289	Sequence 26289, A
6	4094.8	98.3	4543	13	US-10-098-841-173	Sequence 173, App
7	3622.6	87.0	4772	17	US-10-311-034-47	Sequence 47, Appl
8	3174.6	76.2	3276	9	US-09-925-302-242	Sequence 242, App
9	3174.6	76.2	3276	10	US-09-925-302-242	Sequence 242, App
10	962.4	23.1	1768	17	US-10-108-260A-2237	Sequence 2237, Ap
11	454.4	10.9	502	10	US-09-918-995-23795	Sequence 23795, A
c 12	409.6	9.8	505	11	US-09-969-034-4185	Sequence 4185, Ap
13	369	8.9	369	9	US-09-867-701-3627	Sequence 3627, Ap
c 14	320	7.7	396	9	US-09-969-347-122	Sequence 122, App
c 15	320	7.7	396	9	US-09-880-107-1703	Sequence 1703, Ap
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17	291.8	7.0	336	9	US-09-867-701-7412	Sequence 7412, Ap
18	277.8	6.7	599	22	US-10-972-079-12452	Sequence 12452, A
19	273.8	6.6	1747	21	US-10-764-420-2420	Sequence 2420, Ap
20	238.4	5.7	387	17	US-10-264-049-1795	Sequence 1795, Ap
21	228.8	5.5	232	20	US-10-425-115-114670	Sequence 114670,
22	190.8	4.6	337	9	US-09-783-590-2184	Sequence 2184, Ap
23	75.2	1.8	106	17	US-10-242-535A-2596	Sequence 2596, Ap
24	75.2	1.8	106	18	US-10-085-783A-2596	Sequence 2596, Ap
25	65.4	1.6	534	18	US-10-398-877-8	Sequence 8, Appli
26	51.4	1.2	78	9	US-09-783-590-12181	Sequence 12181, A
27	51.2	1.2	3673778	16	US-10-312-841-1	Sequence 1, Appli
28	51	1.2	51	20	US-10-865-478-63	Sequence 63, Appl
29	50.8	1.2	14551	15	US-10-240-485-138	Sequence 138, App
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31	50	1.2	50	17	US-10-131-827-788	Sequence 788, App
c 32	50	1.2	5415	15	US-10-311-455-618	Sequence 618, App
33	49.8	1.2	594	14	US-10-123-155-10	Sequence 10, Appl
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44	49.8	1.2	594	17	US-10-140-864-10	Sequence 10, Appl
45	49.8	1.2	594	18	US-10-142-426-10	Sequence 10, Appl

ALIGNMENTS

RESULT 1
US-09-873-367C-95
; Sequence 95, Application US/09873367C
; Publication No. US20030165839A1
; GENERAL INFORMATION:
; APPLICANT: Young, Paul
; APPLICANT: Soppet, Daniel
; APPLICANT: Endress, Gregory
; APPLICANT: Augustus, Meena
; APPLICANT: Ebner, Reinhard
; APPLICANT: Carter, Kenneth
; TITLE OF INVENTION: Cancer Gene Determination and Therapeutic Screening Using
; TITLE OF INVENTION: Signature Gene Sets
; FILE REFERENCE: 689290-64
; CURRENT APPLICATION NUMBER: US/09/873,367C
; CURRENT FILING DATE: 2003-04-29
; PRIOR APPLICATION NUMBER: U.S. 60/236,891
; PRIOR FILING DATE: 2000-09-29
; PRIOR APPLICATION NUMBER: U.S. 60/236,842
; PRIOR FILING DATE: 2000-09-29
; PRIOR APPLICATION NUMBER: U.S. 60/244,867
; PRIOR FILING DATE: 2000-11-01
; PRIOR APPLICATION NUMBER: U.S. 60/245,084
; PRIOR FILING DATE: 2000-11-01
; NUMBER OF SEQ ID NOS: 1067
; SOFTWARE: PatentIn version 3.0
; SEQ ID NO 95
; LENGTH: 4165
; TYPE: DNA
; ORGANISM: Homo sapiens

US-09-873-367C-95

Query Match 100.0%; Score 4165; DB 10; Length 4165;
Best Local Similarity 100.0%; Pred. No. 0;
Matches 4165; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

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Db       3601 CTTATTAGTTGTGTGGTTGACCCTTGGGGTATACAAATGTCAGTCTGAGTGGTGTCTTAC 3660

Qy       3661 TCCTTTGTTTATAAGTGAATGATTGTGCATGTTTTGTATGTCATAGTATGTCGTCACATA 3720
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       3661 TCCTTTGTTTATAAGTGAATGATTGTGCATGTTTTGTATGTCATAGTATGTCGTCACATA 3720

Qy       3721 AAAGGGAGGGAGCGAAAAACCATTACATTAAGATAATATTGGACCAAACTACTTACTTGC 3780
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Db       3721 AAAGGGAGGGAGCGAAAAACCATTACATTAAGATAATATTGGACCAAACTACTTACTTGC 3780

Qy       3781 TCTAAACAGTTACTTGTACCCCTTAACCTGTCTTCAAAAGTTGCATATAGTTACAGTAGT 3840
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Db       3781 TCTAAACAGTTACTTGTACCCCTTAACCTGTCTTCAAAAGTTGCATATAGTTACAGTAGT 3840

Qy       3841 GTATAAATTAAATATTGTGGAAAAACAGTCTTGTATTTTTCTGTATGTGTGTATATATAT 3900
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Db       3841 GTATAAATTAAATATTGTGGAAAAACAGTCTTGTATTTTTCTGTATGTGTGTATATATAT 3900

Qy       3901 ATAATTATGTACTTCTGGCAATTCTATCTGTATTTAAAGATGTGACAATCTTGACACCAA 3960
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       3901 ATAATTATGTACTTCTGGCAATTCTATCTGTATTTAAAGATGTGACAATCTTGACACCAA 3960

Qy       3961 TTTTAAGAATAGCTGTGAGACCGAATTAAAGATAATCCCTACCAAGTGAAAATTGATGTG 4020
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Db       3961 TTTTAAGAATAGCTGTGAGACCGAATTAAAGATAATCCCTACCAAGTGAAAATTGATGTG 4020

Qy       4021 TGTTAAGAGGGTACAGAATTATCAACTGATTTGGTCAGTTGCTTCCAATGCTGGTTGATT 4080
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Qy       4081 TCCCTCATTGTGTAAACATTGACAGGTATGTGACAAATGGGAAAAAAAATCCAAATAATA 4140
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Qy       4141 AAGTGACATATTGGTGTTCAGCAAT 4165
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Db       4141 AAGTGACATATTGGTGTTCAGCAAT 4165
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RESULT 2
US-10-843-641A-95
; Sequence 95, Application US/10843641A
; Publication No. US20050064454A1
; GENERAL INFORMATION:

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;  APPLICANT: Avalon Pharmaceuticals, Inc.
;  TITLE OF INVENTION: Cancer Gene Determination and Therapeutic Screening Using
;  TITLE OF INVENTION:  Signature Gene Sets
;  FILE REFERENCE: 689290-189
;  CURRENT APPLICATION NUMBER: US/10/843,641A
;  CURRENT FILING DATE:  2004-05-12
;  PRIOR APPLICATION NUMBER: US/09/873,367
;  PRIOR FILING DATE: 2001-06-05
;  PRIOR APPLICATION NUMBER: US/09/954,531
;  PRIOR FILING DATE: 2001-09-18
;  PRIOR APPLICATION NUMBER: US/09/954,456
;  PRIOR FILING DATE: 2001-09-25
;  PRIOR APPLICATION NUMBER: US/09/962,436
;  PRIOR FILING DATE: 2001-09-25
;  PRIOR APPLICATION NUMBER: US/09/962,832
;  PRIOR FILING DATE: 2001-09-25
;  PRIOR APPLICATION NUMBER: US/09/964,824
;  PRIOR FILING DATE: 2001-09-27
;  PRIOR APPLICATION NUMBER: US/09/967,768
;  PRIOR FILING DATE: 2001-09-28
;  PRIOR APPLICATION NUMBER: US/09/968,007
;  PRIOR FILING DATE: 2001-10-02
;  PRIOR APPLICATION NUMBER: US/09/969,347
;  PRIOR FILING DATE: 2001-10-02
;  PRIOR APPLICATION NUMBER: US/09/969,708
;  PRIOR FILING DATE: 2001-10-03
;  Remaining Prior Application data removed - See File Wrapper or PALM.
;  NUMBER OF SEQ ID NOS: 8447
;  SOFTWARE: PatentIn version 3.0
; SEQ ID NO 95
;   LENGTH: 4165
;   TYPE: DNA
;   ORGANISM: Homo sapiens
US-10-843-641A-95

  Query Match             100.0%;  Score 4165;  DB 21;  Length 4165;
  Best Local Similarity   100.0%;  Pred. No. 0;
  Matches 4165;  Conservative    0;  Mismatches    0;  Indels    0;  Gaps    0;

Qy          1 GTCTAAAGAGTGTAAAGACCTAATTACACGGATGCTACAGAGAGATCCCAAGAGAAGGGC 60
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Qy         61 TTCTTTAGAAGAGATTGAAAATCATCCTTGGCTTCAGGGAGTGGACCCTTCACCAGCTAC 120
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Qy        121 AAAGTATAACATTCCCCTTGTGTCATACAAAAATCTCTCGGAAGAGGAGCACAACAGCAT 180
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Qy        181 CATTCAGCGCATGGTGCTTGGGGACATAGCGGATCGAGACGCCATTGTAGAAGCCCTGGA 240
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Qy        241 AACCAACAGGTATAACCATATCACAGCCACATACTTCCTGCTGGCTGAAAGGATCCTGAG 300
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Db        241 AACCAACAGGTATAACCATATCACAGCCACATACTTCCTGCTGGCTGAAAGGATCCTGAG 300

Qy        301 AGAAAAGCAAGAGAAAGAAATACAGACCAGATCTGCAAGCCCGAGCAATATCAAGGCCCA 360
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Qy        361 GTTTAGGCAGTCATGGCCAACCAAAATTGATGTACCCCAGGACCTTGAGGATGACCTCAC 420
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Qy        421 GGCCACTCCTTTGTCCCACGCGACTGTCCCTCAGTCTCCTGCTCGGGCTGCTGACAGTGT 480
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Qy        481 CCTCAATGGCCACAGGAGCAAAGGCCTGTGTGACTCAGCTAAGAAAGATGACCTCCCTGA 540
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Db        481 CCTCAATGGCCACAGGAGCAAAGGCCTGTGTGACTCAGCTAAGAAAGATGACCTCCCTGA 540

Qy        541 GTTGGCTGGACCAGCACTCTCTACGGTGCCACCCGCAAGCTTAAAACCCACAGCCAGTGG 600
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Db        541 GTTGGCTGGACCAGCACTCTCTACGGTGCCACCCGCAAGCTTAAAACCCACAGCCAGTGG 600

Qy        601 GCGGAAGTGTCTGTTCAGGGTGGAAGAAGATGAAGAGGAAGATGAGGAGGACAAGAAACC 660
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Db        601 GCGGAAGTGTCTGTTCAGGGTGGAAGAAGATGAAGAGGAAGATGAGGAGGACAAGAAACC 660

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Qy        721 ATCCAGGAAGAGTGCGCCCGTCCTCAACCAGATCTTTGAGGAAGGGGAATCTGATGATGA 780
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Qy        781 GTTTGACATGGATGAGAATCTGCCTCCCAAGTTGAGCAGGTTAAAGATGAATATAGCTTC 840
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Qy        961 CGGGTTCACCTACTCCTGGCACCGACGGGATAGCAGCGAGGGGCCCCCTGGCAGTGAGGG 1020
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Db        961 CGGGTTCACCTACTCCTGGCACCGACGGGATAGCAGCGAGGGGCCCCCTGGCAGTGAGGG 1020

Qy       1021 GGATGGCGGGGGCCAGAGCAAGCCGAGCAATGCCAGTGGAGGGGTGGACAAGGCCAGCCC 1080
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Db       1021 GGATGGCGGGGGCCAGAGCAAGCCGAGCAATGCCAGTGGAGGGGTGGACAAGGCCAGCCC 1080

Qy       1081 CAGTGAGAACAATGCTGGTGGGGGCAGTCCCTCCAGCGGCTCGGGTGGCAACCCCACCAA 1140
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Db       1081 CAGTGAGAACAATGCTGGTGGGGGCAGTCCCTCCAGCGGCTCGGGTGGCAACCCCACCAA 1140

Qy       1141 TACATCGGGTACCACACGCCGCTGTGCCGGCCCCAGCAACTCCATGCAGCTGGCCTCTCG 1200
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
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Qy       1201 CAGTGCTGGGGAGCTCGTTGAGAGCCTCAAACTCATGAGCCTCTGCCTCGGCTCCCAGCT 1260
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Db       1201 CAGTGCTGGGGAGCTCGTTGAGAGCCTCAAACTCATGAGCCTCTGCCTCGGCTCCCAGCT 1260

Qy       1261 TCATGGGAGCACCAAGTACATTATTGATCCACAGAATGGCTTGTCATTTTCCAGTGTGAA 1320
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Qy       1321 AGTCCAAGAGAAATCTACGTGGAAAATGTGCATTAGCTCCACAGGGAATGCAGGGCAGGT 1380
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Qy       1381 CCCTGCAGTGGGCGGCATAAAGTTTTTCTCTGACCACATGGCAGATACCACCACTGAATT 1440
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Qy       1441 GGAACGGATAAAGAGCAAGAACCTGAAAAATAACGTGCTGCAGCTACCTCTGTGCGAAAA 1500
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Qy       1501 GACCATCTCTGTGAACATCCAGCGGAACCCTAAGGAGGGGCTGCTGTGCGCATCCAGCCC 1560
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Qy      1621 TCAGAGCAGTGAAGACCGGCTCACTTCACTGTTCCATTTGGTTTTACTATTTTAAAGTGG 1680
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Qy      1861 TCTTATAAATGGAGCACTTAGAAATTTGTTGTTCTGCACTTAACCTAGAGAGAGAAAAAA 1920
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Qy      1921 TGCTTTTCTTTGTGAAAAATCTGAATTCCTGTCCTGACCTTCTGTGATGTGGAAACCCTA 1980
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Qy      1981 GGCTCTGAGACACACTCTCTGGTGTCTGAGACAGAACCAAAGCAATAACGTTGTGATGCC 2040
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Qy      2041 CACAGGCCTGGAGCCAGCTAGCGACCTTGTGCCGCCCAGCTGTCCATGGCCCGTGCAGAG 2100
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Qy      2101 CAGAGGACAGTGAGTGTCTGCACTGAGAACCTTAAACCACAGTTGAACATACCCACACCT 2160
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Db      2101 CAGAGGACAGTGAGTGTCTGCACTGAGAACCTTAAACCACAGTTGAACATACCCACACCT 2160

Qy      2161 GTTTGTCTTAAGCTATAGTGTAAAAACAAAGTTTGGGCTCTGAAAATTTAACTGAAAAAG 2220
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db      2161 GTTTGTCTTAAGCTATAGTGTAAAAACAAAGTTTGGGCTCTGAAAATTTAACTGAAAAAG 2220

Qy      2221 ATTTCCTTGTTTTTGTAATAGGTGAGATAAAGTACTTAGATTTATAAGGCAGCTTCCCCT 2280
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Qy      2281 GTAGTGATAAATTACAAGCAGACAATCTTATTTTGTAATGTGATGAAGTGATGATGTCTT 2340
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Qy      2341 AACTCTACTTAGAGAGTGTATGTCTGTCTAACAGAACAAAAAGATGCTCTGTGTAAATTC 2400
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Qy      2401 CTTCCTGTAGGGCACACTGCAGGATTTCCATGTAGATAGAAGAACTATAGGGCCTAGTAC 2460
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Qy      2461 AGAAGGTGCACACAAATGTTGGCAAAGTCAAACCCCATGAATTAAAACCTACTGGAATTT 2520
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Qy      2521 GGTTTTTAGGAGTTTGGTAATTAGATTATCTCTTTTGTTATTTTCATTCAGTTATATCCT 2580
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Qy      2581 TTGGCTCAGCTAGCTTTGAAATTGGCTGATGAAAAAATATACATAAAAGGGTAAAATTCA 2640
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Qy      2641 CACATACAGCAAACAAAAATGCACAAAGCCTGCTTCGTAACTTTTTTTTCTGGAATTGTT 2700
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Qy      2701 TTTCACTTTGCCTTTTTCTGCCAAAACAATAATCAAAGAACTCTTGCTTTAACCTATTCC 2760
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Qy      2881 CTGGGAGTCTTAAAAGCAGTGCTTAGCAACATTGGTGATAGCATGTGGCTGGGACCCAGG 2940
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Qy      3061 TCTTCAAGTACACACTGACTCTGCTACTTTAGGATAAATATATTTTACTCAGAACTCTGA 3120
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Db      3121 ATTTCACAGTATACTTACTAAACTAAGTAAAAATGATACTTAAAATACTTATTTTACTTT 3180

Qy      3181 CTAGACCTAGGCTAGATGTTTTAAGCTACAGCTCTAGTTCATTGTGATATTTATAATTTG 3240
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Db      3181 CTAGACCTAGGCTAGATGTTTTAAGCTACAGCTCTAGTTCATTGTGATATTTATAATTTG 3240

Qy      3241 AAAGCTATGAGAATAGATGTGTGGGTGAAGCCATAGAACATATTTGCTTGAAATTCTTGA 3300
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Db      3241 AAAGCTATGAGAATAGATGTGTGGGTGAAGCCATAGAACATATTTGCTTGAAATTCTTGA 3300

Qy      3301 GCAGGGATCTTATAAAGGGCCAGAAATAAGATGTGTGGTTCACATAGATAGTGAGCGTAA 3360
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Qy      3361 CATCTGTATTAAACATAGGAGAGAAGTTTATAAAGGGCATTGGCAATAAACTCTTTGTTG 3420
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Db      3361 CATCTGTATTAAACATAGGAGAGAAGTTTATAAAGGGCATTGGCAATAAACTCTTTGTTG 3420

Qy      3421 CAGCTGTTTTCCAAGCAGTGTAAATACTTTTTCCTGTGATTATGTATAGCCTTGGAATGG 3480
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Qy      3481 CACCTTTTAACTAACCCATATGTGTTTGGTTTCAATGGTTTTTTATATTCAGATGTATAT 3540
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             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db      3661 TCCTTTGTTTATAAGTGAATGATTGTGCATGTTTTGTATGTCATAGTATGTCGTCACATA 3720

Qy      3721 AAAGGGAGGGAGCGAAAAACCATTACATTAAGATAATATTGGACCAAACTACTTACTTGC 3780
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db      3721 AAAGGGAGGGAGCGAAAAACCATTACATTAAGATAATATTGGACCAAACTACTTACTTGC 3780

Qy      3781 TCTAAACAGTTACTTGTACCCCTTAACCTGTCTTCAAAAGTTGCATATAGTTACAGTAGT 3840
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                ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       3781 TCTAAACAGTTACTTGTACCCCTTAACCTGTCTTCAAAAGTTGCATATAGTTACAGTAGT 3840

Qy       3841 GTATAAATTAAATATTGTGGAAAAACAGTCTTGTATTTTTCTGTATGTGTGTATATATAT 3900
                ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       3841 GTATAAATTAAATATTGTGGAAAAACAGTCTTGTATTTTTCTGTATGTGTGTATATATAT 3900

Qy       3901 ATAATTATGTACTTCTGGCAATTCTATCTGTATTTAAAGATGTGACAATCTTGACACCAA 3960
                ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       3901 ATAATTATGTACTTCTGGCAATTCTATCTGTATTTAAAGATGTGACAATCTTGACACCAA 3960

Qy       3961 TTTTAAGAATAGCTGTGAGACCGAATTAAAGATAATCCCTACCAAGTGAAAATTGATGTG 4020
                ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       3961 TTTTAAGAATAGCTGTGAGACCGAATTAAAGATAATCCCTACCAAGTGAAAATTGATGTG 4020

Qy       4021 TGTTAAGAGGGTACAGAATTATCAACTGATTTGGTCAGTTGCTTCCAATGCTGGTTGATT 4080
                ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       4021 TGTTAAGAGGGTACAGAATTATCAACTGATTTGGTCAGTTGCTTCCAATGCTGGTTGATT 4080

Qy       4081 TCCCTCATTGTGTAAACATTGACAGGTATGTGACAAATGGGAAAAAAAATCCAAATAATA 4140
                ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       4081 TCCCTCATTGTGTAAACATTGACAGGTATGTGACAAATGGGAAAAAAAATCCAAATAATA 4140

Qy       4141 AAGTGACATATTGGTGTTCAGCAAT 4165
                |||||||||||||||||||||||||
Db       4141 AAGTGACATATTGGTGTTCAGCAAT 4165


RESULT 3
US-10-097-340-136
; Sequence 136, Application US/10097340
; Publication No. US20030087250A1
; GENERAL INFORMATION:
;  APPLICANT: John MONAHAN
;  APPLICANT:  Manjula GANNAVARAPU
;  APPLICANT:  Sebastian HOERSCH
;  APPLICANT:  Shubhangi KAMATKAR
;  APPLICANT:  Steve G. KOVATS
;  APPLICANT:  Rachel E. MEYERS
;  APPLICANT:  Michael MORRISEY
;  APPLICANT:  Peter OLANDT
;  APPLICANT:  Ami SEN
;  APPLICANT:  Peter VEIBY
;  APPLICANT:  Gordon B. MILLS
;  APPLICANT:  Robert C. BAST, Jr.
;  APPLICANT:  Karen LU
;  APPLICANT:  Rosemarie SCHMANDT
;  APPLICANT:  Xumei ZHAO
;  APPLICANT:  Karen GLATT
;  TITLE OF INVENTION: Nucleic Acid Molecules and Proteins For The Identification,
;  TITLE OF INVENTION:  Assessment, Prevention, and Therapy of Ovarian Cancer
;  FILE REFERENCE: MRI-030
;  CURRENT APPLICATION NUMBER: US/10/097,340
;  CURRENT FILING DATE:  2002-03-14
;  PRIOR APPLICATION NUMBER: 60/276,025
;  PRIOR FILING DATE: 2001-03-14
;  PRIOR APPLICATION NUMBER: 60/325,149
;  PRIOR FILING DATE: 2001-09-26
;  PRIOR APPLICATION NUMBER: 60/276,026
;  PRIOR FILING DATE: 2001-03-14
;  PRIOR APPLICATION NUMBER: 60/324,967
;  PRIOR FILING DATE: 2001/09/26
;  PRIOR APPLICATION NUMBER: 60/311,732
;  PRIOR FILING DATE: 2001-08-10
;  PRIOR APPLICATION NUMBER: 60/325,102
;  PRIOR FILING DATE: 2001-09-26
;  PRIOR APPLICATION NUMBER: 60/323,580
;  PRIOR FILING DATE: 2001-09-19
;  NUMBER OF SEQ ID NOS: 363
;  SOFTWARE: FastSEQ for Windows Version 4.0
; SEQ ID NO 136
;   LENGTH: 5519
;   TYPE: DNA
;   ORGANISM: Homo sapiens
US-10-097-340-136

  Query Match             100.0%;  Score 4163.4;  DB 14;  Length 5519;
  Best Local Similarity   100.0%;  Pred. No. 0;
  Matches 4164;  Conservative    0;  Mismatches    1;  Indels    0;  Gaps    0;

Qy          1 GTCTAAAGAGTGTAAAGACCTAATTACACGGATGCTACAGAGAGATCCCAAGAGAAGGGC 60
              |||||||||||||||||||| ||||||||||||||||||||||||||||||||||||||
Db       1355 GTCTAAAGAGTGTAAAGACCTAATCACACGGATGCTACAGAGAGATCCCAAGAGAAGGGC 1414

Qy         61 TTCTTTAGAAGAGATTGAAAATCATCCTTGGCTTCAGGGAGTGGACCCTTCACCAGCTAC 120
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       1415 TTCTTTAGAAGAGATTGAAAATCATCCTTGGCTTCAGGGAGTGGACCCTTCACCAGCTAC 1474

Qy        121 AAAGTATAACATTCCCCTTGTGTCATACAAAAATCTCTCGGAAGAGGAGCACAACAGCAT 180
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       1475 AAAGTATAACATTCCCCTTGTGTCATACAAAAATCTCTCGGAAGAGGAGCACAACAGCAT 1534

Qy        181 CATTCAGCGCATGGTGCTTGGGGACATAGCGGATCGAGACGCCATTGTAGAAGCCCTGGA 240
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       1535 CATTCAGCGCATGGTGCTTGGGGACATAGCGGATCGAGACGCCATTGTAGAAGCCCTGGA 1594

Qy        241 AACCAACAGGTATAACCATATCACAGCCACATACTTCCTGCTGGCTGAAAGGATCCTGAG 300
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       1595 AACCAACAGGTATAACCATATCACAGCCACATACTTCCTGCTGGCTGAAAGGATCCTGAG 1654

Qy        301 AGAAAAGCAAGAGAAAGAAATACAGACCAGATCTGCAAGCCCGAGCAATATCAAGGCCCA 360
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       1655 AGAAAAGCAAGAGAAAGAAATACAGACCAGATCTGCAAGCCCGAGCAATATCAAGGCCCA 1714

Qy        361 GTTTAGGCAGTCATGGCCAACCAAAATTGATGTACCCCAGGACCTTGAGGATGACCTCAC 420
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       1715 GTTTAGGCAGTCATGGCCAACCAAAATTGATGTACCCCAGGACCTTGAGGATGACCTCAC 1774

Qy        421 GGCCACTCCTTTGTCCCACGCGACTGTCCCTCAGTCTCCTGCTCGGGCTGCTGACAGTGT 480
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       1775 GGCCACTCCTTTGTCCCACGCGACTGTCCCTCAGTCTCCTGCTCGGGCTGCTGACAGTGT 1834

Qy        481 CCTCAATGGCCACAGGAGCAAAGGCCTGTGTGACTCAGCTAAGAAAGATGACCTCCCTGA 540
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       1835 CCTCAATGGCCACAGGAGCAAAGGCCTGTGTGACTCAGCTAAGAAAGATGACCTCCCTGA 1894

Qy        541 GTTGGCTGGACCAGCACTCTCTACGGTGCCACCCGCAAGCTTAAAACCCACAGCCAGTGG 600
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       1895 GTTGGCTGGACCAGCACTCTCTACGGTGCCACCCGCAAGCTTAAAACCCACAGCCAGTGG 1954

Qy        601 GCGGAAGTGTCTGTTCAGGGTGGAAGAAGATGAAGAGGAAGATGAGGAGGACAAGAAACC 660
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       1955 GCGGAAGTGTCTGTTCAGGGTGGAAGAAGATGAAGAGGAAGATGAGGAGGACAAGAAACC 2014

Qy        661 CATGTCCCTCTCAACACAAGTGGTTTTGCGCCGGAAGCCATCTGTAACCAACCGCCTGAC 720
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       2015 CATGTCCCTCTCAACACAAGTGGTTTTGCGCCGGAAGCCATCTGTAACCAACCGCCTGAC 2074

Qy        721 ATCCAGGAAGAGTGCGCCCGTCCTCAACCAGATCTTTGAGGAAGGGGAATCTGATGATGA 780
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       2075 ATCCAGGAAGAGTGCGCCCGTCCTCAACCAGATCTTTGAGGAAGGGGAATCTGATGATGA 2134

Qy        781 GTTTGACATGGATGAGAATCTGCCTCCCAAGTTGAGCAGGTTAAAGATGAATATAGCTTC 840
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       2135 GTTTGACATGGATGAGAATCTGCCTCCCAAGTTGAGCAGGTTAAAGATGAATATAGCTTC 2194

Qy        841 TCCAGGTACAGTTCACAAACGCTACCACCGGAGGAAAAGTCAGGGCCGGGGCTCCAGCTG 900
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       2195 TCCAGGTACAGTTCACAAACGCTACCACCGGAGGAAAAGTCAGGGCCGGGGCTCCAGCTG 2254

Qy        901 CAGTAGTTCGGAGACCAGTGATGATGATTCTGAAAGCCGGCGGCGGCTCGATAAAGATAG 960
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       2255 CAGTAGTTCGGAGACCAGTGATGATGATTCTGAAAGCCGGCGGCGGCTCGATAAAGATAG 2314

Qy        961 CGGGTTCACCTACTCCTGGCACCGACGGGATAGCAGCGAGGGGCCCCCTGGCAGTGAGGG 1020
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
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Db      2315 CGGGTTCACCTACTCCTGGCACCGACGGGATAGCAGCGAGGGGCCCCCTGGCAGTGAGGG 2374

Qy      1021 GGATGGCGGGGGCCAGAGCAAGCCGAGCAATGCCAGTGGAGGGGTGGACAAGGCCAGCCC 1080
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db      2375 GGATGGCGGGGGCCAGAGCAAGCCGAGCAATGCCAGTGGAGGGGTGGACAAGGCCAGCCC 2434

Qy      1081 CAGTGAGAACAATGCTGGTGGGGGCAGTCCCTCCAGCGGCTCGGGTGGCAACCCCACCAA 1140
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db      2435 CAGTGAGAACAATGCTGGTGGGGGCAGTCCCTCCAGCGGCTCGGGTGGCAACCCCACCAA 2494

Qy      1141 TACATCGGGTACCACACGCCGCTGTGCCGGCCCCAGCAACTCCATGCAGCTGGCCTCTCG 1200
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db      2495 TACATCGGGTACCACACGCCGCTGTGCCGGCCCCAGCAACTCCATGCAGCTGGCCTCTCG 2554

Qy      1201 CAGTGCTGGGGAGCTCGTTGAGAGCCTCAAACTCATGAGCCTCTGCCTCGGCTCCCAGCT 1260
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db      2555 CAGTGCTGGGGAGCTCGTTGAGAGCCTCAAACTCATGAGCCTCTGCCTCGGCTCCCAGCT 2614

Qy      1261 TCATGGGAGCACCAAGTACATTATTGATCCACAGAATGGCTTGTCATTTTCCAGTGTGAA 1320
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db      2615 TCATGGGAGCACCAAGTACATTATTGATCCACAGAATGGCTTGTCATTTTCCAGTGTGAA 2674

Qy      1321 AGTCCAAGAGAAATCTACGTGGAAAATGTGCATTAGCTCCACAGGGAATGCAGGGCAGGT 1380
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db      2675 AGTCCAAGAGAAATCTACGTGGAAAATGTGCATTAGCTCCACAGGGAATGCAGGGCAGGT 2734

Qy      1381 CCCTGCAGTGGGCGGCATAAAGTTTTTCTCTGACCACATGGCAGATACCACCACTGAATT 1440
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db      2735 CCCTGCAGTGGGCGGCATAAAGTTTTTCTCTGACCACATGGCAGATACCACCACTGAATT 2794

Qy      1441 GGAACGGATAAAGAGCAAGAACCTGAAAAATAACGTGCTGCAGCTACCTCTGTGCGAAAA 1500
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db      2795 GGAACGGATAAAGAGCAAGAACCTGAAAAATAACGTGCTGCAGCTACCTCTGTGCGAAAA 2854

Qy      1501 GACCATCTCTGTGAACATCCAGCGGAACCCTAAGGAGGGGCTGCTGTGCGCATCCAGCCC 1560
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db      2855 GACCATCTCTGTGAACATCCAGCGGAACCCTAAGGAGGGGCTGCTGTGCGCATCCAGCCC 2914

Qy      1561 AGCCAGCTGTTGCCATGTCATCTGACTGTGGCCCCATCTGGCCGCTAGCACGCTTCCTGC 1620
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db      2915 AGCCAGCTGTTGCCATGTCATCTGACTGTGGCCCCATCTGGCCGCTAGCACGCTTCCTGC 2974

Qy      1621 TCAGAGCAGTGAAGACCGGCTCACTTCACTGTTCCATTTGGTTTTACTATTTTAAAGTGG 1680
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db      2975 TCAGAGCAGTGAAGACCGGCTCACTTCACTGTTCCATTTGGTTTTACTATTTTAAAGTGG 3034

Qy      1681 GCGTTAGGAGCAATTATTTATTACCTTTCCATTTGTTCGCCTGATGATGTGACAATGCAT 1740
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db      3035 GCGTTAGGAGCAATTATTTATTACCTTTCCATTTGTTCGCCTGATGATGTGACAATGCAT 3094

Qy      1741 GGTCTTTGTGCATGCTGCTAGACACTTTTCTTTCCCAGCCGAAAAGCCTATTATGTAATT 1800
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db      3095 GGTCTTTGTGCATGCTGCTAGACACTTTTCTTTCCCAGCCGAAAAGCCTATTATGTAATT 3154

Qy      1801 TTTACATTCATAATTTTAATGTGGATGATCAGGATTAAATCAAGATATATATCTGGAACC 1860
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db      3155 TTTACATTCATAATTTTAATGTGGATGATCAGGATTAAATCAAGATATATATCTGGAACC 3214

Qy      1861 TCTTATAAATGGAGCACTTAGAAATTTGTTGTTCTGCACTTAACCTAGAGAGAGAAAAAA 1920
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db      3215 TCTTATAAATGGAGCACTTAGAAATTTGTTGTTCTGCACTTAACCTAGAGAGAGAAAAAA 3274

Qy      1921 TGCTTTTCTTTGTGAAAAATCTGAATTCCTGTCCTGACCTTCTGTGATGTGGAAACCCTA 1980
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db      3275 TGCTTTTCTTTGTGAAAAATCTGAATTCCTGTCCTGACCTTCTGTGATGTGGAAACCCTA 3334

Qy      1981 GGCTCTGAGACACACTCTCTGGTGTCTGAGACAGAACCAAAGCAATAACGTTGTGATGCC 2040
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db      3335 GGCTCTGAGACACACTCTCTGGTGTCTGAGACAGAACCAAAGCAATAACGTTGTGATGCC 3394

Qy      2041 CACAGGCCTGGAGCCAGCTAGCGACCTTGTGCCGCCCAGCTGTCCATGGCCCGTGCAGAG 2100
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db      3395 CACAGGCCTGGAGCCAGCTAGCGACCTTGTGCCGCCCAGCTGTCCATGGCCCGTGCAGAG 3454
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Qy      2101 CAGAGGACAGTGAGTGTCTGCACTGAGAACCTTAAACCACAGTTGAACATACCCACACCT 2160
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db      3455 CAGAGGACAGTGAGTGTCTGCACTGAGAACCTTAAACCACAGTTGAACATACCCACACCT 3514

Qy      2161 GTTTGTCTTAAGCTATAGTGTAAAAACAAAGTTTGGGCTCTGAAAATTTAACTGAAAAAG 2220
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db      3515 GTTTGTCTTAAGCTATAGTGTAAAAACAAAGTTTGGGCTCTGAAAATTTAACTGAAAAAG 3574

Qy      2221 ATTTCCTTGTTTTTGTAATAGGTGAGATAAAGTACTTAGATTTATAAGGCAGCTTCCCCT 2280
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db      3575 ATTTCCTTGTTTTTGTAATAGGTGAGATAAAGTACTTAGATTTATAAGGCAGCTTCCCCT 3634

Qy      2281 GTAGTGATAAATTACAAGCAGACAATGTTATTTTGTAATGTGATGAAGTGATGATGTCTT 2340
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db      3635 GTAGTGATAAATTACAAGCAGACAATCTTATTTTGTAATGTGATGAAGTGATGATGTCTT 3694

Qy      2341 AACTCTACTTAGAGAGTGTATGTCTGTCTAACAGAACAAAAAGATGCTCTGTGTAAATTC 2400
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db      3695 AACTCTACTTAGAGAGTGTATGTCTGTCTAACAGAACAAAAAGATGCTCTGTGTAAATTC 3754

Qy      2401 CTTCCTGTAGGGCACACTGCAGGATTTCCATGTAGATAGAAGAACTATAGGGCCTAGTAC 2460
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db      3755 CTTCCTGTAGGGCACACTGCAGGATTTCCATGTAGATAGAAGAACTATAGGGCCTAGTAC 3814

Qy      2461 AGAAGGTGCACACAAATGTTGGCAAAGTCAAACCCCATGAATTAAAACCTACTGGAATTT 2520
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db      3815 AGAAGGTGCACACAAATGTTGGCAAAGTCAAACCCCATGAATTAAAACCTACTGGAATTT 3874

Qy      2521 GGTTTTTAGGAGTTTGGTAATTAGATTATCTCTTTTGTTATTTTCATTCAGTTATATCCT 2580
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db      3875 GGTTTTTAGGAGTTTGGTAATTAGATTATCTCTTTTGTTATTTTCATTCAGTTATATCCT 3934

Qy      2581 TTGGCTCAGCTAGCTTTGAAATTGGCTGATGAAAAAATATACATAAAAGGGTAAAATTCA 2640
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db      3935 TTGGCTCAGCTAGCTTTGAAATTGGCTGATGAAAAAATATACATAAAAGGGTAAAATTCA 3994

Qy      2641 CACATACAGCAAACAAAAATGCACAAAGCCTGCTTCGTAACTTTTTTTTCTGGAATTGTT 2700
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db      3995 CACATACAGCAAACAAAAATGCACAAAGCCTGCTTCGTAACTTTTTTTTCTGGAATTGTT 4054

Qy      2701 TTTCACTTTGCCTTTTTCTGCCAAAACAATAATCAAAGAACTCTTGCTTTAACCTATTCC 2760
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db      4055 TTTCACTTTGCCTTTTTCTGCCAAAACAATAATCAAAGAACTCTTGCTTTAACCTATTCC 4114

Qy      2761 TGTACAAAGACTGTTTTTGACCAGATAATCATCTGTTGTGGCATTCTATCTTGTAGGACA 2820
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db      4115 TGTACAAAGACTGTTTTTGACCAGATAATCATCTGTTGTGGCATTCTATCTTGTAGGACA 4174

Qy      2821 CTGTATATTGCAAATTGCTGATTATGGAAGGGGCCAGTTGCTGTTTTTTTCATGCAGTGCC 2880
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db      4175 CTGTATATTGCAAATTGCTGATTATGGAAGGGGCCAGTTGCTGTTTTTTTCATGCAGTGCC 4234

Qy      2881 CTGGGAGTCTTAAAAGCAGTGCTTAGCAACATTGGTGATAGCATGTGGCTGGGACCCAGG 2940
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db      4235 CTGGGAGTCTTAAAAGCAGTGCTTAGCAACATTGGTGATAGCATGTGGCTGGGACCCAGG 4294

Qy      2941 GCCCTTCCCCACTCTTCAGCCCCGAGTCATGTGTCTGAGGTGACGGACTGAGACGCATCT 3000
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db      4295 GCCCTTCCCCACTCTTCAGCCCCGAGTCATGTGTCTGAGGTGACGGACTGAGACGCATCT 4354

Qy      3001 GGTCCTGTAATTCAGAGAGTGGGCACATCACCAAAGAACTGCATTGCTGTGGTCACTGTT 3060
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db      4355 GGTCCTGTAATTCAGAGAGTGGGCACATCACCAAAGAACTGCATTGCTGTGGTCACTGTT 4414

Qy      3061 TCTTCAAGTACACACTGACTCTGCTACTTTAGGATAAATATATTTTACTCAGAACTCTGA 3120
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db      4415 TCTTCAAGTACACACTGACTCTGCTACTTTAGGATAAATATATTTTACTCAGAACTCTGA 4474

Qy      3121 ATTTCACAGTATACTTACTAAACTAAGTAAAAATGATACTTAAAATACTTATTTTACTTT 3180
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db      4475 ATTTCACAGTATACTTACTAAACTAAGTAAAAATGATACTTAAAATACTTATTTTACTTT 4534
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Qy       3181 CTAGACCTAGGCTAGATGTTTTAAGCTACAGCTCTAGTTCATTGTGATATTTATAATTTG 3240
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       4535 CTAGACCTAGGCTAGATGTTTTAAGCTACAGCTCTAGTTCATTGTGATATTTATAATTTG 4594

Qy       3241 AAAGCTATGAGAATAGATGTGTGGGTGAAGCCATAGAACATATTTGCTTGAAATTCTTGA 3300
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       4595 AAAGCTATGAGAATAGATGTGTGGGTGAAGCCATAGAACATATTTGCTTGAAATTCTTGA 4654

Qy       3301 GCAGGGATCTTATAAAGGGCCAGAAATAAGATGTGTGGTTCACATAGATAGTGAGCGTAA 3360
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       4655 GCAGGGATCTTATAAAGGGCCAGAAATAAGATGTGTGGTTCACATAGATAGTGAGCGTAA 4714

Qy       3361 CATCTGTATTAAACATAGGAGAGAAGTTTATAAAGGGCATTGGCAATAAACTCTTTGTTG 3420
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       4715 CATCTGTATTAAACATAGGAGAGAAGTTTATAAAGGGCATTGGCAATAAACTCTTTGTTG 4774

Qy       3421 CAGCTGTTTTCCAAGCAGTGTAAATACTTTTTCCTGTGATTATGTATAGCCTTGGAATGG 3480
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       4775 CAGCTGTTTTCCAAGCAGTGTAAATACTTTTTCCTGTGATTATGTATAGCCTTGGAATGG 4834

Qy       3481 CACCTTTTAACTAACCCATATGTGTTTGGTTTCAATGGTTTTTTATATTCAGATGTATAT 3540
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       4835 CACCTTTTAACTAACCCATATGTGTTTGGTTTCAATGGTTTTTTATATTCAGATGTATAT 4894

Qy       3541 ATGGTGCTCACTTTAGGATCAGCAGTGTTGACCATTTATGCTGCATAGCTGTATTATAGC 3600
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       4895 ATGGTGCTCACTTTAGGATCAGCAGTGTTGACCATTTATGCTGCATAGCTGTATTATAGC 4954

Qy       3601 CTTATTAGTTGTGTGGTTGACCCTTGGGGTATACAAATGTCAGTCTGAGTGGTGTCTTAC 3660
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       4955 CTTATTAGTTGTGTGGTTGACCCTTGGGGTATACAAATGTCAGTCTGAGTGGTGTCTTAC 5014

Qy       3661 TCCTTTGTTTATAAGTGAATGATTGTGCATGTTTTGTATGTCATAGTATGTCGTCACATA 3720
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       5015 TCCTTTGTTTATAAGTGAATGATTGTGCATGTTTTGTATGTCATAGTATGTCGTCACATA 5074

Qy       3721 AAAGGGAGGGAGCGAAAAACCATTACATTAAGATAATATTGGACCAAACTACTTACTTGC 3780
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       5075 AAAGGGAGGGAGCGAAAAACCATTACATTAAGATAATATTGGACCAAACTACTTACTTGC 5134

Qy       3781 TCTAAACAGTTACTTGTACCCCTTAACCTGTCTTCAAAAGTTGCATATAGTTACAGTAGT 3840
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       5135 TCTAAACAGTTACTTGTACCCCTTAACCTGTCTTCAAAAGTTGCATATAGTTACAGTAGT 5194

Qy       3841 GTATAAATTAAATATTGTGGAAAAACAGTCTTGTATTTTTCTGTATGTGTGTATATATAT 3900
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       5195 GTATAAATTAAATATTGTGGAAAAACAGTCTTGTATTTTTCTGTATGTGTGTATATATAT 5254

Qy       3901 ATAATTATGTACTTCTGGCAATTCTATCTGTATTTAAAGATGTGACAATCTTGACACCAA 3960
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       5255 ATAATTATGTACTTCTGGCAATTCTATCTGTATTTAAAGATGTGACAATCTTGACACCAA 5314

Qy       3961 TTTTAAGAATAGCTGTGAGACCGAATTAAAGATAATCCCTACCAAGTGAAAATTGATGTG 4020
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       5315 TTTTAAGAATAGCTGTGAGACCGAATTAAAGATAATCCCTACCAAGTGAAAATTGATGTG 5374

Qy       4021 TGTTAAGAGGGTACAGAATTATCAACTGATTTGGTCAGTTGCTTCCAATGCTGGTTGATT 4080
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       5375 TGTTAAGAGGGTACAGAATTATCAACTGATTTGGTCAGTTGCTTCCAATGCTGGTTGATT 5434

Qy       4081 TCCCTCATTGTGTAAACATTGACAGGTATGTGACAAATGGGAAAAAAAATCCAAATAATA 4140
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       5435 TCCCTCATTGTGTAAACATTGACAGGTATGTGACAAATGGGAAAAAAAATCCAAATAATA 5494

Qy       4141 AAGTGACATATTGGTGTTCAGCAAT 4165
              |||||||||||||||||||||||||
Db       5495 AAGTGACATATTGGTGTTCAGCAAT 5519
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RESULT 4
US-10-354-358-69
; Sequence 69, Application US/10354358
; Publication No. US20030157082A1

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; GENERAL INFORMATION:
;  APPLICANT: Millennium Pharmaceuticals, Inc
;  APPLICANT:  Hunter, John Joseph
;  APPLICANT:  MacBeth, Kyle J.
;  APPLICANT:  Tsai, Fong-Ying
;  APPLICANT:  Lesoon, Andrea
;  APPLICANT:  Lightcap, Eric S.
;  APPLICANT:  Williamson, Mark
;  APPLICANT:  Rudolph-Owen, Laura A.
;  TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR TREATING
;  TITLE OF INVENTION:  CANCER USING 140, 1470, 1686, 2089, 2427, 3702, 5891, 6428,
;  TITLE OF INVENTION:  7181, 7660, 25641, 69583, 49863, 8897, 1682, 17667, 9235,
;  TITLE OF INVENTION:  3703, 14171, 10359, 1660, 1450, 18894, 2088, 32427, 2160,
;  TITLE OF INVENTION:  9252, 9389, 1642, 85269, 10297, 1584, 9525, 14124, 4469,
;  TITLE OF INVENTION:  8990, 2100, 9288, 64698, 10480, 20893, 33230, 1586, 9943,
;  TITLE OF INVENTION:  16334, 68862, 9011, 14031, 6178, 21225, 1420, 32236, 2099,
;  TITLE OF INVENTION:  2150, 26583, 2784, 8941, 9811, 27444, 50566 OR 66428 MOLECULES
;  FILE REFERENCE: MPI02-020P1RNOMNIM
;  CURRENT APPLICATION NUMBER: US/10/354,358
;  CURRENT FILING DATE:  2003-01-30
;  PRIOR APPLICATION NUMBER: US 60/353,600
;  PRIOR FILING DATE: 2002-01-31
;  PRIOR APPLICATION NUMBER: US 60/364,517
;  PRIOR FILING DATE: 2002-03-15
;  PRIOR APPLICATION NUMBER: US 60/371,075
;  PRIOR FILING DATE: 2002-04-09
;  PRIOR APPLICATION NUMBER: US 60/371,507
;  PRIOR FILING DATE: 2002-04-10
;  PRIOR APPLICATION NUMBER: US 60/372,984
;  PRIOR FILING DATE: 2002-04-16
;  PRIOR APPLICATION NUMBER: US 60/374,194
;  PRIOR FILING DATE: 2002-04-19
;  PRIOR APPLICATION NUMBER: US 60/382,995
;  PRIOR FILING DATE: 2002-05-24
;  PRIOR APPLICATION NUMBER: US 60/385,023
;  PRIOR FILING DATE: 2002-05-31
;  PRIOR APPLICATION NUMBER: US 60/388,853
;  PRIOR FILING DATE: 2002-06-14
;  PRIOR APPLICATION NUMBER: US 60/389,395
;  PRIOR FILING DATE: 2002-06-17
;  Remaining Prior Application data removed - See File Wrapper or PALM.
;  NUMBER OF SEQ ID NOS: 122
;  SOFTWARE: FastSEQ for Windows Version 4.0
; SEQ ID NO 69
;    LENGTH: 5519
;    TYPE: DNA
;    ORGANISM: Homo sapiens
;    FEATURE:
;    NAME/KEY: CDS
;    LOCATION: (642)...(2939)
US-10-354-358-69

  Query Match            100.0%;  Score 4163.4;  DB 16;  Length 5519;
  Best Local Similarity   100.0%;  Pred. No. 0;
  Matches 4164;  Conservative    0;  Mismatches    1;  Indels    0;  Gaps    0;

Qy          1 GTCTAAAGAGTGTAAAGACCTAATTACACGGATGCTACAGAGAGATCCCAAGAGAAGGGC 60
              |||||||||||||||||||||| ||||||||||||||||||||||||||||||||||||
Db       1355 GTCTAAAGAGTGTAAAGACCTAATCACACGGATGCTACAGAGAGATCCCAAGAGAAGGGC 1414

Qy         61 TTCTTTAGAAGAGATTGAAAATCATCCTTGGCTTCAGGGAGTGGACCCTTCACCAGCTAC 120
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       1415 TTCTTTAGAAGAGATTGAAAATCATCCTTGGCTTCAGGGAGTGGACCCTTCACCAGCTAC 1474

Qy        121 AAAGTATAACATTCCCCTTGTGTCATACAAAAATCTCTCGGAAGAGGAGCACAACAGCAT 180
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       1475 AAAGTATAACATTCCCCTTGTGTCATACAAAAATCTCTCGGAAGAGGAGCACAACAGCAT 1534

Qy        181 CATTCAGCGCATGGTGCTTGGGGACATAGCGGATCGAGACGCCATTGTAGAAGCCCTGGA 240
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       1535 CATTCAGCGCATGGTGCTTGGGGACATAGCGGATCGAGACGCCATTGTAGAAGCCCTGGA 1594

Qy        241 AACCAACAGGTATAACCATATCACAGCCACATACTTCCTGCTGGCTGAAAGGATCCTGAG 300
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Db      1595 AACCAACAGGTATAACCATATCACAGCCACATACTTCCTGCTGGCTGAAAGGATCCTGAG 1654
Qy       301 AGAAAAGCAAGAGAAAGAAATACAGACCAGATCTGCAAGCCCGAGCAATATCAAGGCCCA 360
Db      1655 AGAAAAGCAAGAGAAAGAAATACAGACCAGATCTGCAAGCCCGAGCAATATCAAGGCCCA 1714
Qy       361 GTTTAGGCAGTCATGGCCAACCAAAATTGATGTACCCCAGGACCTTGAGGATGACCTCAC 420
Db      1715 GTTTAGGCAGTCATGGCCAACCAAAATTGATGTACCCCAGGACCTTGAGGATGACCTCAC 1774
Qy       421 GGCCACTCCTTTGTCCCACGCGACTGTCCCTCAGTCTCCTGCTCGGGCTGCTGACAGTGT 480
Db      1775 GGCCACTCCTTTGTCCCACGCGACTGTCCCTCAGTCTCCTGCTCGGGCTGCTGACAGTGT 1834
Qy       481 CCTCAATGGCCACAGGAGCAAAGGCCTGTGTGACTCAGCTAAGAAAGATGACCTCCCTGA 540
Db      1835 CCTCAATGGCCACAGGAGCAAAGGCCTGTGTGACTCAGCTAAGAAAGATGACCTCCCTGA 1894
Qy       541 GTTGGCTGGACCAGCACTCTCTACGGTGCCACCCGCAAGCTTAAAACCCACAGCCAGTGG 600
Db      1895 GTTGGCTGGACCAGCACTCTCTACGGTGCCACCCGCAAGCTTAAAACCCACAGCCAGTGG 1954
Qy       601 GCGGAAGTGTCTGTTCAGGGTGGAAGAAGATGAAGAGGAAGATGAGGAGGACAAGAAACC 660
Db      1955 GCGGAAGTGTCTGTTCAGGGTGGAAGAAGATGAAGAGGAAGATGAGGAGGACAAGAAACC 2014
Qy       661 CATGTCCCTCTCAACACAAGTGGTTTTGCGCCGGAAGCCATCTGTAACCAACCGCCTGAC 720
Db      2015 CATGTCCCTCTCAACACAAGTGGTTTTGCGCCGGAAGCCATCTGTAACCAACCGCCTGAC 2074
Qy       721 ATCCAGGAAGAGTGCGCCCGTCCTCAACCAGATCTTTGAGGAAGGGGAATCTGATGATGA 780
Db      2075 ATCCAGGAAGAGTGCGCCCGTCCTCAACCAGATCTTTGAGGAAGGGGAATCTGATGATGA 2134
Qy       781 GTTTGACATGGATGAGAATCTGCCTCCCAAGTTGAGCAGGTTAAAGATGAATATAGCTTC 840
Db      2135 GTTTGACATGGATGAGAATCTGCCTCCCAAGTTGAGCAGGTTAAAGATGAATATAGCTTC 2194
Qy       841 TCCAGGTACAGTTCACAAACGCTACCACCGGAGGAAAAGTCAGGGCCGGGGCTCCAGCTG 900
Db      2195 TCCAGGTACAGTTCACAAACGCTACCACCGGAGGAAAAGTCAGGGCCGGGGCTCCAGCTG 2254
Qy       901 CAGTAGTTCGGAGACCAGTGATGATGATTCTGAAAGCCGGCGGCGGCTCGATAAAGATAG 960
Db      2255 CAGTAGTTCGGAGACCAGTGATGATGATTCTGAAAGCCGGCGGCGGCTCGATAAAGATAG 2314
Qy       961 CGGGTTCACCTACTCCTGGCACCGACGGGATAGCAGCGAGGGGCCCCCTGGCAGTGAGGG 1020
Db      2315 CGGGTTCACCTACTCCTGGCACCGACGGGATAGCAGCGAGGGGCCCCCTGGCAGTGAGGG 2374
Qy      1021 GGATGGCGGGGGCCAGAGCAAGCCGAGCAATGCCAGTGGAGGGGTGGACAAGGCCAGCCC 1080
Db      2375 GGATGGCGGGGGCCAGAGCAAGCCGAGCAATGCCAGTGGAGGGGTGGACAAGGCCAGCCC 2434
Qy      1081 CAGTGAGAACAATGCTGGTGGGGGCAGTCCCTCCAGCGGCTCGGGTGGCAACCCCACCAA 1140
Db      2435 CAGTGAGAACAATGCTGGTGGGGGCAGTCCCTCCAGCGGCTCGGGTGGCAACCCCACCAA 2494
Qy      1141 TACATCGGGTACCACACGCCGCTGTGCCGGCCCCAGCAACTCCATGCAGCTGGCCTCTCG 1200
Db      2495 TACATCGGGTACCACACGCCGCTGTGCCGGCCCCAGCAACTCCATGCAGCTGGCCTCTCG 2554
Qy      1201 CAGTGCTGGGGAGCTCGTTGAGAGCCTCAAACTCATGAGCCTCTGCCTCGGCTCCCAGCT 1260
Db      2555 CAGTGCTGGGGAGCTCGTTGAGAGCCTCAAACTCATGAGCCTCTGCCTCGGCTCCCAGCT 2614
Qy      1261 TCATGGGAGCACCAAGTACATTATTGATCCACAGAATGGCTTGTCATTTTCCAGTGTGAA 1320
Db      2615 TCATGGGAGCACCAAGTACATTATTGATCCACAGAATGGCTTGTCATTTTCCAGTGTGAA 2674
Qy      1321 AGTCCAAGAGAAATCTACGTGGAAAATGTGCATTAGCTCCACAGGGAATGCAGGGCAGGT 1380
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Db      2675 AGTCCAAGAGAAATCTACGTGGAAAATGTGCATTAGCTCCACAGGGAATGCAGGGCAGGT 2734
Qy      1381 CCCTGCAGTGGGCGGCATAAAGTTTTTCTCTGACCACATGGCAGATACCACCACTGAATT 1440
Db      2735 CCCTGCAGTGGGCGGCATAAAGTTTTTCTCTGACCACATGGCAGATACCACCACTGAATT 2794
Qy      1441 GGAACGGATAAAGAGCAAGAACCTGAAAAATAACGTGCTGCAGCTACCTCTGTGCGAAAA 1500
Db      2795 GGAACGGATAAAGAGCAAGAACCTGAAAAATAACGTGCTGCAGCTACCTCTGTGCGAAAA 2854
Qy      1501 GACCATCTCTGTGAACATCCAGCGGAACCCTAAGGAGGGGCTGCTGTGCGCATCCAGCCC 1560
Db      2855 GACCATCTCTGTGAACATCCAGCGGAACCCTAAGGAGGGGCTGCTGTGCGCATCCAGCCC 2914
Qy      1561 AGCCAGCTGTTGCCATGTCATCTGACTGTGGCCCCATCTGGCCGCTAGCACGCTTCCTGC 1620
Db      2915 AGCCAGCTGTTGCCATGTCATCTGACTGTGGCCCCATCTGGCCGCTAGCACGCTTCCTGC 2974
Qy      1621 TCAGAGCAGTGAAGACCGGCTCACTTCACTGTTCCATTTGGTTTTACTATTTTAAAGTGG 1680
Db      2975 TCAGAGCAGTGAAGACCGGCTCACTTCACTGTTCCATTTGGTTTTACTATTTTAAAGTGG 3034
Qy      1681 GCGTTAGGAGCAATTATTTATTACCTTTCCATTTGTTCGCCTGATGATGTGACAATGCAT 1740
Db      3035 GCGTTAGGAGCAATTATTTATTACCTTTCCATTTGTTCGCCTGATGATGTGACAATGCAT 3094
Qy      1741 GGTCTTTGTGCATGCTGCTAGACACTTTTCTTTCCCAGCCGAAAAGCCTATTATGTAATT 1800
Db      3095 GGTCTTTGTGCATGCTGCTAGACACTTTTCTTTCCCAGCCGAAAAGCCTATTATGTAATT 3154
Qy      1801 TTTACATTCATAATTTTAATGTGGATGATCAGGATTAAATCAAGATATATATCTGGAACC 1860
Db      3155 TTTACATTCATAATTTTAATGTGGATGATCAGGATTAAATCAAGATATATATCTGGAACC 3214
Qy      1861 TCTTATAAATGGAGCACTTAGAAATTTGTTGTTCTGCACTTAACCTAGAGAGAGAAAAAA 1920
Db      3215 TCTTATAAATGGAGCACTTAGAAATTTGTTGTTCTGCACTTAACCTAGAGAGAGAAAAAA 3274
Qy      1921 TGCTTTTCTTTGTGAAAAATCTGAATTCCTGTCCTGACCTTCTGTGATGTGGAAACCCTA 1980
Db      3275 TGCTTTTCTTTGTGAAAAATCTGAATTCCTGTCCTGACCTTCTGTGATGTGGAAACCCTA 3334
Qy      1981 GGCTCTGAGACACACTCTCTGGTGTCTGAGACAGAACCAAAGCAATAACGTTGTGATGCC 2040
Db      3335 GGCTCTGAGACACACTCTCTGGTGTCTGAGACAGAACCAAAGCAATAACGTTGTGATGCC 3394
Qy      2041 CACAGGCCTGGAGCCAGCTAGCGACCTTGTGCCGCCCAGCTGTCCATGGCCCGTGCAGAG 2100
Db      3395 CACAGGCCTGGAGCCAGCTAGCGACCTTGTGCCGCCCAGCTGTCCATGGCCCGTGCAGAG 3454
Qy      2101 CAGAGGACAGTGAGTGTCTGCACTGAGAACCTTAAACCACAGTTGAACATACCCACACCT 2160
Db      3455 CAGAGGACAGTGAGTGTCTGCACTGAGAACCTTAAACCACAGTTGAACATACCCACACCT 3514
Qy      2161 GTTTGTCTTAAGCTATAGTGTAAAAACAAAGTTTGGGCTCTGAAAATTTAACTGAAAAAG 2220
Db      3515 GTTTGTCTTAAGCTATAGTGTAAAAACAAAGTTTGGGCTCTGAAAATTTAACTGAAAAAG 3574
Qy      2221 ATTTCCTTGTTTTTGTAATAGGTGAGATAAAGTACTTAGATTTATAAGGCAGCTTCCCCT 2280
Db      3575 ATTTCCTTGTTTTTGTAATAGGTGAGATAAAGTACTTAGATTTATAAGGCAGCTTCCCCT 3634
Qy      2281 GTAGTGATAAATTACAAGCAGACAATCTTATTTTGTAATGTGATGAAGTGATGATGTCTT 2340
Db      3635 GTAGTGATAAATTACAAGCAGACAATCTTATTTTGTAATGTGATGAAGTGATGATGTCTT 3694
Qy      2341 AACTCTACTTAGAGAGTGTATGTCTGTCTAACAGAAGAAAAAGATGCTCTGTGTAAATTC 2400
Db      3695 AACTCTACTTAGAGAGTGTATGTCTGTCTAACAGAACAAAAAGATGCTCTGTGTAAATTC 3754
Qy      2401 CTTCCTGTAGGGCACACTGCAGGATTTCCATGTAGATAGAAGAACTATAGGGCCTAGTAC 2460
Db      3755 CTTCCTGTAGGGCACACTGCAGGATTTCCATGTAGATAGAAGAACTATAGGGCCTAGTAC 3814
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Qy       2461 AGAAGGTGCACACAAATGTTGGCAAAGTCAAACCCCATGAATTAAAACCTACTGGAATTT 2520
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Db       3815 AGAAGGTGCACACAAATGTTGGCAAAGTCAAACCCCATGAATTAAAACCTACTGGAATTT 3874

Qy       2521 GGTTTTTAGGAGTTTGGTAATTAGATTATCTCTTTTGTTATTTTCATTCAGTTATATCCT 2580
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Db       3875 GGTTTTTAGGAGTTTGGTAATTAGATTATCTCTTTTGTTATTTTCATTCAGTTATATCCT 3934

Qy       2581 TTGGCTCAGCTAGCTTTGAAATTGGCTGATGAAAAAATATACATAAAAGGGTAAAATTCA 2640
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       3935 TTGGCTCAGCTAGCTTTGAAATTGGCTGATGAAAAAATATACATAAAAGGGTAAAATTCA 3994

Qy       2641 CACATACAGCAAACAAAAATGCACAAAGCCTGCTTCGTAACTTTTTTTTCTGGAATTGTT 2700
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Db       3995 CACATACAGCAAACAAAAATGCACAAAGCCTGCTTCGTAACTTTTTTTTCTGGAATTGTT 4054

Qy       2701 TTTCACTTTGCCTTTTTCTGCCAAAACAATAATCAAAGAACTCTTGCTTTAACCTATTCC 2760
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       4055 TTTCACTTTGCCTTTTTCTGCCAAAACAATAATCAAAGAACTCTTGCTTTAACCTATTCC 4114

Qy       2761 TGTACAAAGACTGTTTTTGACCAGATAATCATCTGTTGTGGCATTCTATCTTGTAGGACA 2820
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       4115 TGTACAAAGACTGTTTTTGACCAGATAATCATCTGTTGTGGCATTCTATCTTGTAGGACA 4174

Qy       2821 CTGTATATTGCAAATTGCTGATTATGGAAGGGGCCAGTTGCTGTTTTTTCATGCAGTGCC 2880
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Db       4175 CTGTATATTGCAAATTGCTGATTATGGAAGGGGCCAGTTGCTGTTTTTTCATGCAGTGCC 4234

Qy       2881 CTGGGAGTCTTAAAAGCAGTGCTTAGCAACATTGGTGATAGCATGTGGCTGGGACCCAGG 2940
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Db       4235 CTGGGAGTCTTAAAAGCAGTGCTTAGCAACATTGGTGATAGCATGTGGCTGGGACCCAGG 4294

Qy       2941 GCCCTTCCCCACTCTTCAGCCCCGAGTCATGTGTCTGAGGTGACGGACTGAGACGCATCT 3000
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Db       4295 GCCCTTCCCCACTCTTCAGCCCCGAGTCATGTGTCTGAGGTGACGGACTGAGACGCATCT 4354

Qy       3001 GGTCCTGTAATTCAGAGAGTGGGCACATCACCAAAGAACTGCATTGCTGTGGTCACTGTT 3060
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       4355 GGTCCTGTAATTCAGAGAGTGGGCACATCACCAAAGAACTGCATTGCTGTGGTCACTGTT 4414

Qy       3061 TCTTCAAGTACACACTGACTCTGCTACTTTAGGATAAATATATTTTACTCAGAACTCTGA 3120
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Db       4415 TCTTCAAGTACACACTGACTCTGCTACTTTAGGATAAATATATTTTACTCAGAACTCTGA 4474

Qy       3121 ATTTCACAGTATACTTACTAAACTAAGTAAAAATGATACTTAAAATACTTATTTTACTTT 3180
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Db       4475 ATTTCACAGTATACTTACTAAACTAAGTAAAAATGATACTTAAAATACTTATTTTACTTT 4534

Qy       3181 CTAGACCTAGGCTAGATGTTTTAAGCTACAGCTCTAGTTCATTGTGATATTTATAATTTG 3240
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       4535 CTAGACCTAGGCTAGATGTTTTAAGCTACAGCTCTAGTTCATTGTGATATTTATAATTTG 4594

Qy       3241 AAAGCTATGAGAATAGATGTGTGGGTGAAGCCATAGAACATATTTGCTTGAAATTCTTGA 3300
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Db       4595 AAAGCTATGAGAATAGATGTGTGGGTGAAGCCATAGAACATATTTGCTTGAAATTCTTGA 4654

Qy       3301 GCAGGGATCTTATAAAGGGCCAGAAATAAGATGTGTGGTTCACATAGATAGTGAGCGTAA 3360
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       4655 GCAGGGATCTTATAAAGGGCCAGAAATAAGATGTGTGGTTCACATAGATAGTGAGCGTAA 4714

Qy       3361 CATCTGTATTAAACATAGGAGAGAAGTTTATAAAGGGCATTGGCAATAAACTCTTTGTTG 3420
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       4715 CATCTGTATTAAACATAGGAGAGAAGTTTATAAAGGGCATTGGCAATAAACTCTTTGTTG 4774

Qy       3421 CAGCTGTTTTCCAAGCAGTGTAAATACTTTTTCCTGTGATTATGTATAGCCTTGGAATGG 3480
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       4775 CAGCTGTTTTCCAAGCAGTGTAAATACTTTTTCCTGTGATTATGTATAGCCTTGGAATGG 4834

Qy       3481 CACCTTTTAACTAACCCATATGTGTTTGGTTTCAATGGTTTTTTATATTCAGATGTATAT 3540
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       4835 CACCTTTTAACTAACCCATATGTGTTTGGTTTCAATGGTTTTTTATATTCAGATGTATAT 4894

Qy       3541 ATGGTGCTCACTTTAGGATCAGCAGTGTTGACCATTTATGCTGCATAGCTGTATTATAGC 3600
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       4895 ATGGTGCTCACTTTAGGATCAGCAGTGTTGACCATTTATGCTGCATAGCTGTATTATAGC 4954

Qy       3601 CTTATTAGTTGTGTGGTTGACCCTTGGGGTATACAAATGTCAGTCTGAGTGGTGTCTTAC 3660
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       4955 CTTATTAGTTGTGTGGTTGACCCTTGGGGTATACAAATGTCAGTCTGAGTGGTGTCTTAC 5014

Qy       3661 TCCTTTGTTTATAAGTGAATGATTGTGCATGTTTTGTATGTCATAGTATGTCGTCACATA 3720
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       5015 TCCTTTGTTTATAAGTGAATGATTGTGCATGTTTTGTATGTCATAGTATGTCGTCACATA 5074

Qy       3721 AAAGGGAGGGAGCGAAAAACCATTACATTAAGATAATATTGGACCAAACTACTTACTTGC 3780
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       5075 AAAGGGAGGGAGCGAAAAACCATTACATTAAGATAATATTGGACCAAACTACTTACTTGC 5134

Qy       3781 TCTAAACAGTTACTTGTACCCCTTAACCTGTCTTCAAAAGTTGCATATAGTTACAGTAGT 3840
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       5135 TCTAAACAGTTACTTGTACCCCTTAACCTGTCTTCAAAAGTTGCATATAGTTACAGTAGT 5194

Qy       3841 GTATAAATTAAATATTGTGGAAAAACAGTCTTGTATTTTTCTGTATGTGTGTATATATAT 3900
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       5195 GTATAAATTAAATATTGTGGAAAAACAGTCTTGTATTTTTCTGTATGTGTGTATATATAT 5254

Qy       3901 ATAATTATGTACTTCTGGCAATTCTATCTGTATTTAAAGATGTGACAATCTTGACACCAA 3960
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Db       5255 ATAATTATGTACTTCTGGCAATTCTATCTGTATTTAAAGATGTGACAATCTTGACACCAA 5314

Qy       3961 TTTTAAGAATAGCTGTGAGACCGAATTAAAGATAATCCCTACCAAGTGAAAATTGATGTG 4020
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Db       5315 TTTTAAGAATAGCTGTGAGACCGAATTAAAGATAATCCCTACCAAGTGAAAATTGATGTG 5374

Qy       4021 TGTTAAGAGGGTACAGAATTATCAACTGATTTGGTCAGTTGCTTCCAATGCTGGTTGATT 4080
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Db       5375 TGTTAAGAGGGTACAGAATTATCAACTGATTTGGTCAGTTGCTTCCAATGCTGGTTGATT 5434

Qy       4081 TCCCTCATTGTGTAAACATTGACAGGTATGTGACAAATGGGAAAAAAAATCCAAATAATA 4140
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Db       5435 TCCCTCATTGTGTAAACATTGACAGGTATGTGACAAATGGGAAAAAAAATCCAAATAATA 5494

Qy       4141 AAGTGACATATTGGTGTTCAGCAAT 4165
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Db       5495 AAGTGACATATTGGTGTTCAGCAAT 5519
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RESULT 5
US-10-425-114-26289
; Sequence 26289, Application US/10425114
; Publication No. US20040034888A1
; GENERAL INFORMATION:
; APPLICANT: Liu, Jingdong
; APPLICANT: Zhou, Yihua
; APPLICANT: Kovalic, David K.
; APPLICANT: Screen, Steven E
; APPLICANT: Tabaska, Jack E
; APPLICANT: Cao, Yongwei
; TITLE OF INVENTION: Nucleic Acid Molecules and Other Molecules Associated With
; TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement
; FILE REFERENCE: 38-21(53313)B
; CURRENT APPLICATION NUMBER: US/10/425,114
; CURRENT FILING DATE: 2003-04-28
; NUMBER OF SEQ ID NOS: 73128
; SEQ ID NO 26289
; LENGTH: 4176
; TYPE: DNA
; ORGANISM: Homo sapiens
; FEATURE:
; OTHER INFORMATION: Clone ID: LIB4119-206-H10_FLI
US-10-425-114-26289

Query Match 99.4%; Score 4138.2; DB 18; Length 4176;
Best Local Similarity 99.9%; Pred. No. 0;
Matches 4162; Conservative 0; Mismatches 3; Indels 2; Gaps 2;

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Qy       1 GTCTAAAGAGTGTAAAGACCTAATTACACGGATGCTACAGAGAGATCCCAAGAGAAGGGC 60
Db       8 GTCTAAAGAGTGTAAAGACCTAATCACACGGATGCTACAGAGAGATCCCAAGAGAAGGGC 67

Qy      61 TTCTTTAGAAGAGATTGAAAATCATCCTTGGCTTCAGGGAGTGGACCCTTCACCAGCTAC 120
Db      68 TTCTTTAGAAGAGATTGAAAATCATCCTTGGCTTCAGGGAGTGGACCCTTCACCAGCTAC 127

Qy     121 AAAGTATAACATTCCCCTTGTGTCATACAAAAATCTCTCGGAAGAGGAGCACAACAGCAT 180
Db     128 AAAGTATAACATTCCCCTTGTGTCATACAAAAATCTCTCGGAAGAGGAGCACAACAGCAT 187

Qy     181 CATTCAGCGCATGGTGCTTGGGGACATAGCGGATCGAGACGCCATTGTAGAAGCCCTGGA 240
Db     188 CATTCAGCGCATGGTGCTTGGGGACATAGCGGATCGAGACGCCATTGTAGAAGCCCTGGA 247

Qy     241 AACCAACAGGTATAACCATATCACAGCCACATACTTCCTGCTGGCTGAAAGGATCCTGAG 300
Db     248 AACCAACAGGTATAACCATATCACAGCCACATACTTCCTGCTGGCTGAAAGGATCCTGAG 307

Qy     301 AGAAAAGCAAGAGAAAGAAATACAGACCAGATCTGCAAGCCCGAGCAATATCAAGGCCCA 360
Db     308 AGAAAAGCAAGAGAAAGAAATACAGACCAGATCTGCAAGCCCGAGCAATATCAAGGCCCA 367

Qy     361 GTTTAGGCAGTCATGGCCAACCAAAATTGATGTACCCCAGGACCTTGAGGATGACCTCAC 420
Db     368 GTTTAGGCAGTCATGGCCAACCAAAATTGATGTACCCCAGGACCTTGAGGATGACCTCAC 427

Qy     421 GGCCACTCCTTTGTCCCACGCGACTGTCCCTCAGTCTCCTGCTCGGGCTGCTGACAGTGT 480
Db     428 GGCCACTCCTTTGTCCCACGCGACTGTCCCTCAGTCTCCTGCTCGGGCTGCTGACAGTGT 487

Qy     481 CCTCAATGGCCACAGGAGCAAAGGCCTGTGTGACTCAGCTAAGAAAGATGACCTCCCTGA 540
Db     488 CCTCAATGGCCACAGGAGCAAAGGCCTGTGTGACTCAGCTAAGAAAGATGACCTCCCTGA 547

Qy     541 GTTGGCTGGACCAGCACTCTCTACGGTGCCACCCGCAAGCTTAAAACCCACAGCCAGTGG 600
Db     548 GTTGGCTGGACCAGCACTCTCTACGGTGCCACCCGCAAGCTTAAAACCCACAGCCAGTGG 607

Qy     601 GCGGAAGTGTCTGTTCAGGGTGGAAGAAGATGAAGAGGAAGATGAGGAGGACAAGAAACC 660
Db     608 GCGGAAGTGTCTGTTCAGGGTGGAAGAAGATGAAGAGGAAGATGAGGAGGACAAGAAACC 667

Qy     661 CATGTCCCTCTCAACACAAGTGGTTTTGCGCCGGAAGCCATCTGTAACCAACCGCCTGAC 720
Db     668 CATGTCCCTCTCAACACAAGTGGTTTTGCGCCGGAAGCCATCTGTAACCAACCGCCTGAC 727

Qy     721 ATCCAGGAAGAGTGCGCCCGTCCTCAACCAGATCTTTGAGGAAGGGGAATCTGATGATGA 780
Db     728 ATCCAGGAAGAGTGCGCCCGTCCTCAACCAGATCTTTGAGGAAGGGGAATCTGACGATGA 787

Qy     781 GTTTGACATGGATGAGAATCTGCCTCCCAAGTTGAGCAGGTTAAAGATGAATATAGCTTC 840
Db     788 GTTTGACATGGATGAGAATCTGCCTCCCAAGTTGAGCAGGTTAAAGATGAATATAGCTTC 847

Qy     841 TCCAGGTACAGTTCACAAACGCTACCACCGGAGGAAAAGTCAGGGCCGGGGCTCCAGCTG 900
Db     848 TCCAGGTACAGTTCACAAACGCTACCACCGGAGGAAAAGTCAGGGCCGGGGCTCCAGCTG 907

Qy     901 CAGTAGTTCGGAGACCAGTGATGATGATTCTGAAAGCCGGCGGCGGCTCGATAAAGATAG 960
Db     908 CAGTAGTTCGGAGACCAGTGATGATGATTCTGAAAGCCGGCGGCGGCTCGATAAAGATAG 967

Qy     961 CGGGTTCACCTACTCCTGGCACCGACGGGATAGCAGCGAGGGGCCCCCTGGCAGTGAGGG 1020
Db     968 CGGGTTCACCTACTCCTGGCACCGACGGGATAGCAGCGAGGGGCCCCCTGGCAGTGAGGG 1027

Qy    1021 GGATGGCGGGGGCCAGAGCAAGCCGAGCAATGCCAGTGGAGGGGTGGACAAGGCCAGCCC 1080
Db    1028 GGATGGCGGGGGCCAGAGCAAGCCAAGCAATGCCAGTGGAGGGGTGGACAAGGCCAGCCC 1087

Qy    1081 CAGTGAGAACAATGCTGGTGGGGGCAGTCCCTCCAGCGGCTCGGGTGGCAACCCCACCAA 1140
Db    1088 CAGTGAGAACAATGCTGGTGGGGGCAGTCCCTCCAGCGGCTCGGGTGGCAACCCCACCAA 1147

Qy    1141 TACATCGGGTACCACACGCCGCTGTGCCGGCCCCAGCAACTCCATGCAGCTGGCCTCTCG 1200
Db    1148 TACATCGGGTACCACACGCCGCTGTGCCGGCCCCAGCAACTCCATGCAGCTGGCCTCTCG 1207

Qy    1201 CAGTGCTGGGGAGCTCGTTGAGAGCCTCAAACTCATGAGCCTCTGCCTCGGCTCCCAGCT 1260
Db    1208 CAGTGCTGGGGAGCTCGTTGAGAGCCTCAAACTCATGAGCCTCTGCCTCGGCTCCCAGCT 1267

Qy    1261 TCATGGGAGCACCAAGTACATTATTGATCCACAGAATGGCTTGTCATTTTCCAGTGTGAA 1320
Db    1268 TCATGGGAGCACCAAGTACATTATTGATCCACAGAATGGCTTGTCATTTTCCAGTGTGAA 1327

Qy    1321 AGTCCAAGAGAAATCTACGTGGAAAATGTGCATTAGCTCCACAGGGAATGCAGGGCAGGT 1380
Db    1328 AGTCCAAGAGAAATCTACGTGGAAAATGTGCATTAGCTCCACAGGGAATGCAGGGCAGGT 1387

Qy    1381 CCCTGCAGTGGGCGGCATAAAGTTTTTCTCTGACCACATGGCAGATACCACCACTGAATT 1440
Db    1388 CCCTGCAGTGGGCGGCATAAAGTTTTTCTCTGACCACATGGCAGATACCACCACTGAATT 1447

Qy    1441 GGAACGGATAAAGAGCAAGAACCTGAAAAATAACGTGCTGCAGCTACCTCTGTGCGAAAA 1500
Db    1448 GGAACGGATAAAGAGCAAGAACCTGAAAAATAACGTGCTGCAGCTACCTCTGTGCGAAAA 1507

Qy    1501 GACCATCTCTGTGAACATCCAGCGGAACCCTAAGGAGGGGCTGCTGTGCGCATCCAGCCC 1560
Db    1508 GACCATCTCTGTGAACATCCAGCGGAACCCTAAGGAGGGGCTGCTGTGCGCATCCAGCCC 1567

Qy    1561 AGCCAGCTGTTGCCATGTCATCTGACTGTGGCCCCATCTGGCCGCTAGCACGCTTCCTGC 1620
Db    1568 AGCCAGCTGTTGCCATGTCATCTGACTGTGGCCCCATCTGGCCGCTAGCACGCTTCCTGC 1627

Qy    1621 TCAGAGCAGTGAAGACCGGCTCACTTCACTGTTCCATTTGGTTTTACTATTTTAAAGTGG 1680
Db    1628 TCAGAGCAGTGAAGACCGGCTCACTTCACTGTTCCATTTGGTTTTACTATTTTAAAGTGG 1687

Qy    1681 GCGTTAGGAGCAATTATTTATTACCTTTCCATTTGTTCGCCTGATGATGTGACAATGCAT 1740
Db    1688 GCGTTAGGAGCAATTATTTATTACCTTTCCATTTGTTCGCCTGATGATGTGACAATGCAT 1747

Qy    1741 GGTCTTTGTGCATGCTGCTAGACACTTTTCTTTCCCAGCCGAAAAGCCTATTATGTAATT 1800
Db    1748 GGTCTTTGTGCATGCTGCTAGACACTTTTCTTTCCCAGCCGAAAAGCCTATTATGTAATT 1807

Qy    1801 TTTACATTCATAATTTTAATGTGGATGATCAGGATTAAATCAAGATATATATCTGGAACC 1860
Db    1808 TTTACATTCATAATTTTAATGTGGATGATCAGGATTAAATCAAGATATATATCTGGAACC 1867

Qy    1861 TCTTATAAATGGAGCACTTAGAAATTTGTTGTTCTGCACTTAACCTAGAGAGAGAAAAAA 1920
Db    1868 TCTTATAAATGGAGCACTTAGAAATTTGTTGTTCTGCACTTAACCTAGAGAGAGAAAAAA 1927

Qy    1921 TGC-TTTTCTTTGTGAAAAATCTGAATTCCTGTCCTGACCTTCTGTGATGTGGAAACCCT 1979
Db    1928 TGCTTTTTCTTTGTGAAAAATCTGAATTCCTGTCCTGACCTTCTGTGATGTGGAAACCCT 1987

Qy    1980 AGGCTCTGAGACACACTCTCTGGTGTCTGAGACAGAACCAAAGCAATAACGTTGTGATGC 2039
Db    1988 AGGCTCTGAGACACACTCTCTGGTGTCTGAGACAGAACCAAAGCAATAACGTTGTGATGC 2047

Qy    2040 CCACAGGCCTGGAGCCAGCTAGCGACCTTGTGCCGCCCAGCTGTCCATGGCCCGTGCAGA 2099
Db    2048 CCACAGGCCTGGAGCCAGCTAGCGACCTTGTGCCGCCCAGCTGTCCATGGCCCGTGCAGA 2107

Qy    2100 GCAGAGGACAGTGAGTGTCTGCACTGAGAACCTTAAACCACAGTTGAACATACCCACACC 2159
Db    2108 GCAGAGGACAGTGAGTGTCTGCACTGAGAACCTTAAACCACAGTTGAACATACCCACACC 2167

Qy    2160 TGTTTGTCTTAAGCTATAGTGTAAAAACAAAGTTTGGGCTCTGAAAATTTAACTGAAAAA 2219
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Db      2168 TGTTTGTCTTAAGCTATAGTGTAAAAACAAAGTTTGGGCTCTGAAAATTTAACTGAAAAA 2227
Qy      2220 GATTTCCTTGTTTTTGTAATAGGTGAGATAAAGTACTTAGATTTATAAGGCAGCTTCCCC 2279
Db      2228 GATTTCCTTGTTTTTGTAATAGGTGAGATAAAGTACTTAGATTTATAAGGCAGCTTCCCC 2287
Qy      2280 TGTAGTGATAAATTACAAGCAGACAATCTTATTTTGTAATGTGATGAAGTGATGATGTCT 2339
Db      2288 TGTAGTGATAAATTACAAGCAGACAATCTTATTTTGTAATGTGATGAAGTGATGATGTCT 2347
Qy      2340 TAACTCTACTTAGAGAGTGTATGTCTGTCTAACAGAACAAAAAGATGCTCTGTGTAAATT 2399
Db      2348 TAACTCTACTTAGAGAGTGTATGTCTGTCTAACAGAACAAAAAGATGCTCTGTGTAAATT 2407
Qy      2400 CCTTCCTGTAGGGCACACTGCAGGATTTCCATGTAGATAGAAGAACTATAGGGCCTAGTA 2459
Db      2408 CCTTCCTGTAGGGCACACTGCAGGATTTCCATGTAGATAGAAGAACTATAGGGCCTAGTA 2467
Qy      2460 CAGAAGGTGCACACAAATGTTGGCAAAGTC-AAACCCCATGAATTAAAACCTACTGGAAT 2518
Db      2468 CAGAAGGTGCACACAAATGTTGGCAAAGTCAAAACCCCATGAATTAAAACCTACTGGAAT 2527
Qy      2519 TTGGTTTTTAGGAGTTTGGTAATTAGATTATCTCTTTTGTTATTTTCATTCAGTTATATC 2578
Db      2528 TTGGTTTTTAGGAGTTTGGTAATTAGATTATCTCTTTTGTTATTTTCATTCAGTTATATC 2587
Qy      2579 CTTTGGCTCAGCTAGCTTTGAAATTGGCTGATGAAAAAATATACATAAAAGGGTAAAATT 2638
Db      2588 CTTTGGCTCAGCTAGCTTTGAAATTGGCTGATGAAAAAATATACATAAAAGGGTAAAATT 2647
Qy      2639 CACACATACAGCAAACAAAAATGCACAAAGCCTGCTTCGTAACTTTTTTTTCTGGAATTG 2698
Db      2648 CACACATACAGCAAACAAAAATGCACAAAGCCTGCTTCGTAACTTTTTTTTCTGGAATTG 2707
Qy      2699 TTTTTCACTTTGCCTTTTTCTGCCAAAACAATAATCAAAGAACTCTTGCTTTAACCTATT 2758
Db      2708 TTTTTCACTTTGCCTTTTTCTGCCAAAACAATAATCAAAGAACTCTTGCTTTAACCTATT 2767
Qy      2759 CCTGTACAAAGACTGTTTTTGACCAGATAATCATCTGTTGTGGCATTCTATCTTGTAGGA 2818
Db      2768 CCTGTACAAAGACTGTTTTTGACCAGATAATCATCTGTTGTGGCATTCTATCTTGTAGGA 2827
Qy      2819 CACTGTATATTGCAAATTGCTGATTATGGAAGGGGCCAGTTGCTGTTTTTTCATGCAGTG 2878
Db      2828 CACTGTATATTGCAAATTGCTGATTATGGAAGGGGCCAGTTGCTGTTTTTTCATGCAGTG 2887
Qy      2879 CCCTGGGAGTCTTAAAAGCAGTGCTTAGCAACATTGGTGATAGCATGTGGCTGGGACCCA 2938
Db      2888 CCCTGGGAGTCTTAAAAGCAGTGCTTAGCAACATTGGTGATAGCATGTGGCTGGGACCCA 2947
Qy      2939 GGGCCCTTCCCCACTCTTCAGCCCCGAGTCATGTGTCTGAGGTGACGGACTGAGACGCAT 2998
Db      2948 GGGCCCTTCCCCACTCTTCAGCCCCGAGTCATGTGTCTGAGGTGACGGACTGAGACGCAT 3007
Qy      2999 CTGGTCCTGTAATTCAGAGAGTGGGCACATCACCAAAGAACTGCATTGCTGTGGTCACTG 3058
Db      3008 CTGGTCCTGTAATTCAGAGAGTGGGCACATCACCAAAGAACTGCATTGCTGTGGTCACTG 3067
Qy      3059 TTTCTTCAAGTACACACTGACTCTGCTACTTTAGGATAAATATATTTTACTCAGAACTCT 3118
Db      3068 TTTCTTCAAGTACACACTGACTCTGCTACTTTAGGATAAATATATTTTACTCAGAACTCT 3127
Qy      3119 GAATTTCACAGTATACTTACTAAACTAAGTAAAAATGATACTTAAAATACTTATTTTACT 3178
Db      3128 GAATTTCACAGTATACTTACTAAACTAAGTAAAAATGATACTTAAAATACTTATTTTACT 3187
Qy      3179 TTCTAGACCTAGGCTAGATGTTTTAAGCTACAGCTCTAGTTCATTGTGATATTTATAATT 3238
Db      3188 TTCTAGACCTAGGCTAGATGTTTTAAGCTACAGCTCTAGTTCATTGTGATATTTATAATT 3247
Qy      3239 TGAAAGCTATGAGAATAGATGTGTGGGTGAAGCCATAGAACATATTTGCTTGAAATTCTT 3298
Db      3248 TGAAAGCTATGAGAATAGATGTGTGGGTGAAGCCATAGAACATATTTGCTTGAAATTCTT 3307
Qy      3299 GAGCAGGGATCTTATAAAGGGCCAGAAATAAGATGTGTGGTTCACATAGATAGTGAGCGT 3358
Db      3308 GAGCAGGGATCTTATAAAGGGCCAGAAATAAGATGTGTGGTTCACATAGATAGTGAGCGT 3367
Qy      3359 AACATCTGTATTAAACATAGGAGAGAAGTTTATAAAGGGCATTGGCAATAAACTCTTTGT 3418
Db      3368 AACATCTGTATTAAACATAGGAGAGAAGTTTATAAAGGGCATTGGCAATAAACTCTTTGT 3427
Qy      3419 TGCAGCTGTTTTCCAAGCAGTGTAAATACTTTTTCCTGTGATTATGTATAGCCTTGGAAT 3478
Db      3428 TGCAGCTGTTTTCCAAGCAGTGTAAATACTTTTTCCTGTGATTATGTATAGCCTTGGAAT 3487
Qy      3479 GGCACCTTTTAACTAACCCATATGTGTTTGGTTTCAATGGTTTTTTATATTCAGATGTAT 3538
Db      3488 GGCACCTTTTAACTAACCCATATGTGTTTGGTTTCAATGGTTTTTTATATTCAGATGTAT 3547
Qy      3539 ATATGGTGCTCACTTTAGGATCAGCAGTGTTGACCATTTATGCTGCATAGCTGTATTATA 3598
Db      3548 ATATGGTGCTCACTTTAGGATCAGCAGTGTTGACCATTTATGCTGCATAGCTGTATTATA 3607
Qy      3599 GCCTTATTAGTTGTGTGGTTGACCCTTGGGGTATACAAATGTCAGTCTGAGTGGTGTCTT 3658
Db      3608 GCCTTATTAGTTGTGTGGTTGACCCTTGGGGTATACAAATGTCAGTCTGAGTGGTGTCTT 3667
Qy      3659 ACTCCTTTGTTTATAAGTGAATGATTGTGCATGTTTTGTATGTCATAGTATGTCGTCACA 3718
Db      3668 ACTCCTTTGTTTATAAGTGAATGATTGTGCATGTTTTGTATGTCATAGTATGTCGTCACA 3727
Qy      3719 TAAAAGGGAGGGAGCGAAAAACCATTACATTAAGATAATATTGGACCAAACTACTTACTT 3778
Db      3728 TAAAAGGGAGGGAGCGAAAAACCATTACATTAAGATAATATTGGACCAAACTACTTACTT 3787
Qy      3779 GCTCTAAACAGTTACTTGTACCCCTTAACCTGTCTTCAAAAGTTGCATATAGTTACAGTA 3838
Db      3788 GCTCTAAACAGTTACTTGTACCCCTTAACCTGTCTTCAAAAGTTGCATATAGTTACAGTA 3847
Qy      3839 GTGTATAAATTAAATATTGTGGAAAAACAGTCTTGTATTTTTCTGTATGTGTGTATATAT 3898
Db      3848 GTGTATAAATTAAATATTGTGGAAAAACAGTCTTGTATTTTTCTGTATGTGTGTATATAT 3907
Qy      3899 ATATAATTATGTACTTCTGGCAATTCTATCTGTATTTAAAGATGTGACAATCTTGACACC 3958
Db      3908 ATATAATTATGTACTTCTGGCAATTCTATCTGTATTTAAAGATGTGACAATCTTGACACC 3967
Qy      3959 AATTTTAAGAATAGCTGTGAGACCGAATTAAAGATAATCCCTACCAAGTGAAAATTGATG 4018
Db      3968 AATTTTAAGAATAGCTGTGAGACCGAATTAAAGATAATCCCTACCAAGTGAAAATTGATG 4027
Qy      4019 TGTGTTAAGAGGGTACAGAATTATCAACTGATTTGGTCAGTTGCTTCCAATGCTGGTTGA 4078
Db      4028 TGTGTTAAGAGGGTACAGAATTATCAACTGATTTGGTCAGTTGCTTCCAATGCTGGTTGA 4087
Qy      4079 TTTCCCTCATTGTGTAAACATTGACAGGTATGTGACAAATGGGAAAAAAAATCCAAATAA 4138
Db      4088 TTTCCCTCATTGTGTAAACATTGACAGGTATGTGACAAATGGGAAAAAAAATCCAAATAA 4147
Qy      4139 TAAAGTGACATATTGGTGTTCAGCAAT 4165
Db      4148 TAAAGTGACATATTGGTGTTCAGCAAT 4174
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RESULT 6
US-10-098-841-173
; Sequence 173, Application US/10098841
; Publication No. US20020197679A1
; GENERAL INFORMATION:
; APPLICANT: Tang, Y. Tom
; APPLICANT: Liu, Chenghua
; APPLICANT: Asundi, Vinod
; APPLICANT: Xu, Chongjun
; APPLICANT: Zhou, Ping

; APPLICANT: Ma, Yunqing
; APPLICANT: Wang, Jian-Rui
; APPLICANT: Zhao, Qing A.
; APPLICANT: Ren, Feiyan
; APPLICANT: Chen, Rui-hong
; APPLICANT: Wang, Dunrui
; APPLICANT: Wang, Zhiwei
; APPLICANT: Wehrman, Tom
; APPLICANT: Zhang, Jie
; APPLICANT: Qian, Xiaohong B.
; APPLICANT: Drmanac, Radoje T.
; TITLE OF INVENTION: No. US20020197679A1el Nucleic Acids and
; TITLE OF INVENTION: Polypeptides
; FILE REFERENCE: 784CIP2
; CURRENT APPLICATION NUMBER: US/10/098,841
; CURRENT FILING DATE: 2002-03-13
; PRIOR APPLICATION NUMBER: 09/598,042
; PRIOR FILING DATE: 2000-06-20
; PRIOR APPLICATION NUMBER: 09/552,317
; PRIOR FILING DATE: 2000-04-25
; PRIOR APPLICATION NUMBER: 09/488,725
; PRIOR FILING DATE: 2000-01-21
; NUMBER OF SEQ ID NOS: 331
; SOFTWARE: pt_FL_genes Version 1.0
; SEQ ID NO 173
; LENGTH: 4543
; TYPE: DNA
; ORGANISM: Homo sapiens
; FEATURE:
; NAME/KEY: CDS
; LOCATION: (266)..(1945)
US-10-098-841-173

Query Match 98.3%; Score 4094.8; DB 13; Length 4543;
Best Local Similarity 99.8%; Pred. No. 0;
Matches 4163; Conservative 0; Mismatches 2; Indels 7; Gaps 6;

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Qy          1 GTCTAAAGAGTGTAAAGACCTAATTACACGGATGCTACAGAGAGATCCCAAGAGAAGGGC 60
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Db        361 GTCTAAAGAGTGTAAAGACCTAATCACACGGATGCTACAGAGAGATCCCAAGAGAAGGGC 420

Qy         61 TTCTTTAGAAGAGATTGAAAATCATCCTTGGCTTCAGGGAGTGGACCCTTCACCAGCTAC 120
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        421 TTCTTTAGAAGAGATTGAAAATCATCCTTGGCTTCAGGGAGTGGACCCTTCACCAGCTAC 480

Qy        121 AAAGTATAACATTCCCCTTGTGTCATACAAAAATCTCTCGGAAGAGGAGCACAACAGCAT 180
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Db        481 AAAGTATAACATTCCCCTTGTGTCATACAAAAATCTCTCGGAAGAGGAGCACAACAGCAT 540

Qy        181 CATTCAGCGCATGGTGCTTGGGGACATAGCGGATCGAGACGCCATTGTAGAAGCCCTGGA 240
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Db        541 CATTCAGCGCATGGTGCTTGGGGACATAGCGGATCGAGACGCCATTGTAGAAGCCCTGGA 600

Qy        241 AACCAACAGGTATAACCATATCACAGCCACATACTTCCTGCTGGCTGAAAGGATCCTGAG 300
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Db        601 AACCAACAGGTATAACCATATCACAGCCACATACTTCCTGCTGGCTGAAAGGATCCTGAG 660

Qy        301 AGAAAAGCAAGAGAAAGAAATACAGACCAGATCTGCAAGCCCGAGCAATATCAAGGCCCA 360
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Db        661 AGAAAAGCAAGAGAAAGAAATACAGACCAGATCTGCAAGCCCGAGCAATATCAAGGCCCA 720

Qy        361 GTTTAGGCAGTCATGGCCAACCAAAATTGATGTACCCCAGGACCTTGAGGATGACCTCAC 420
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Db        721 GTTTAGGCAGTCATGGCCAACCAAAATTGATGTACCCCAGGACCTTGAGGATGACCTCAC 780

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Qy        481 CCTCAATGGCCACAGGAGCAAAGGCCTGTGTGACTCAGCTAAGAAAGATGACCTCCCTGA 540
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Db        841 CCTCAATGGCCACAGGAGCAAAGGCCTGTGTGACTCAGCTAAGAAAGATGACCTCCCTGA 900

Qy        541 GTTGGCTGGACCAGCACTCTCTACGGTGCCACCCGCAAGCTTAAAACCCACAGCCAGTGG 600
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Qy        601 GCGGAAGTGTCTGTTCAGGGTGGAAGAAGATGAAGAGGAAGATGAGGAGGACAAGAAACC 660
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Qy        661 CATGTCCCTCTCAACACAAGTGGTTTTGCGCCGGAAGCCATCTGTAACCAACCGCCTGAC 720
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Db       1021 CATGTCCCTCTCAACACAAGTGGTTTTGCGCCGGAAGCCATCTGTAACCAACCGCCTGAC 1080

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Qy        781 GTTTGACATGGATGAGAATCTGCCTCCCAAGTTGAGCAGGTTAAAGATGAATATAGCTTC 840
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Qy        841 TCCAGGTACAGTTCACAAACGCTACCACCGGAGGAAAAGTCAGGGCCGGGGCTCCAGCTG 900
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Qy        901 CAGTAGTTCGGAGACCAGTGATGATGATTCTGAAAGCCGGCGGCGGCTCGATAAAGATAG 960
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Db       1261 CAGTAGTTCGGAGACCAGTGATGATGATTCTGAAAGCCGGCGGCGGCTCGATAAAGATAG 1320

Qy        961 CGGGTTCACCTACTCCTGGCACCGACGGGATAGCAGCGAGGGGCCCCCTGGCAGTGAGGG 1020
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Db       1321 CGGGTTCACCTACTCCTGGCACCGACGGGATAGCAGCGAGGGGCCCCCTGGCAGTGAGGG 1380

Qy       1021 GGATGGCGGGGGCCAGAGCAAGCCGAGCAATGCCAGTGGAGGGGTGGACAAGGCCAGCCC 1080
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Qy       1081 CAGTGAGAACAATGCTGGTGGGGGCAGTCCCTCCAGCGGCTCGGGTGGCAACCCCACCAA 1140
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Db       1441 CAGTGAGAACAATGCTGGTGGGGGCAGTCCCTCCAGCGGCTCGGGTGGCAACCCCACCAA 1500

Qy       1141 TACATCGGGTACCACACGCCGCTGTGCCGGCCCCAGCAACTCCATGCAGCTGGCCTCTCG 1200
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Db       1501 TACATCGGGTACCACACGCCGCTGTGCCGGCCCCAGCAACTCCATGCAGCTGGCCTCTCG 1560

Qy       1201 CAGTGCTGGGGAGCTCGTTGAGAGCCTCAAACTCATGAGCCTCTGCCTCGGCTCCCAGCT 1260
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Db       1561 CAGTGCTGGGGAGCTCGTTGAGAGCCTCAAACTCATGAGCCTCTGCCTCGGCTCCCAGCT 1620

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Db       1621 TCATGGGAGCACCAAGTACATTATTGATCCACAGAATGGCTTGTCATTTTCCAGTGTGAA 1680

Qy       1321 AGTCCAAGAGAAATCTACGTGGAAAATGTGCATTAGCTCCACAGGGAATGCAGGGCAGGT 1380
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Db       1681 AGTCCAAGAGAAATCTACGTGGAAAATGTGCATTAGCTCCACAGGGAATGCAGGGCAGGT 1740

Qy       1381 CCCTGCAGTGGGCGGCATAAAGTTTTTCTCTGACCACATGGCAGATACCACCACTGAATT 1440
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Db       1741 CCCTGCAGTGGGCGGCATAAAGTTTTTCTCTGACCACATGGCAGATACCACCACTGAATT 1800

Qy       1441 GGAACGGATAAAGAGCAAGAACCTGAAAAATAACGTGCTGCAGCTACCTCTGTGCGAAAA 1500
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       1801 GGAACGGATAAAGAGCAAGAACCTGAAAAATAACGTGCTGCAGCTACCTCTGTGCGAAAA 1860

Qy       1501 GACCATCTCTGTGAACATCCAGCGGAACCCTAAGGAGGGGCTGCTGTGCGCATCCAGCCC 1560
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       1861 GACCATCTCTGTGAACATCCAGCGGAACCCTAAGGAGGGGCTGCTGTGCGCATCCAGCCC 1920

Qy       1561 AGCCAGCTGTTGCCATGTCATCTGACTGTGGCCCCATCTGGCCGCTAGCACGCTTCCTGC 1620
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       1921 AGCCAGCTGTTGCCATGTCATCTGACTGTGGCCCCATCTGGCCGCTAGCACGCTTCCTGC 1980

Qy       1621 TCAGAGCAGTGAAGACCGGCTCACTTCACTGTTCCATTTGGTTTTACTATTTTAAAGTGG 1680
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Db      1981 TCAGAGCAGTGAAGACCGGCTCACTTCACTGTTCCATTTGGTTTTACTATTTTAAAGTGG 2040
Qy      1681 GCGTTAGGAGCAATTATTTATTACCTTTCCATTTGTTCGCCTGATGATGTGACAATGCAT 1740
Db      2041 GCGTTAGGAGCAATTATTTATTACCTTTCCATTTGTTCGCCTGATGATGTGACAATGCAT 2100
Qy      1741 GGTCTTTGTGCATGCTGCTAGACACTTTTCTTTCCCAGCCGAAAAGCCTATTATGTAATT 1800
Db      2101 GGTCTTTGTGCATGCTGCTAGACACTTTTCTTTCCCAGCCGAAAAGCCTATTATGTAATT 2160
Qy      1801 TTTACATTCATAATTTTAATGTGGATGATCAGGATTAAATCAAGATATATATCTGGAACC 1860
Db      2161 TTTACATTCATAATTTTAATGTGGATGATCAGGATTAAATCAAGATATATATCTGGAACC 2220
Qy      1861 TCTTATAAATGGAGCACTTAGAAATTTGTTGTTCTGCACTTAACCTAGAGAGAGAAAAAA 1920
Db      2221 TCTTATAAATGGAGCACTTAGAAATTTGTTGTTCTGCACTTAACCTAGAGAGAGAAAAAA 2280
Qy      1921 TGC-TTTTCTTTGTGAAAAATCTGAATTCCTGTCCTGACCTTCTGTGATGTGGAAACCCT 1979
Db      2281 TGCTTTTTCTTTGTGAAAAATCTGAATTCCTGTCCTGACCTTCTGTGATGTGGAAACCCT 2340
Qy      1980 AGGCTCTGAGACACACTCTCTGGTGTCTGAGACAGAACCAAAGCAATAACGTTGTGATGC 2039
Db      2341 AGGCTCTGAGACACACTCTCTGGTGTCTGAGACAGAACCAAAGCAATAACGTTGTGATGC 2400
Qy      2040 CCACAGGCCTGGAGCCAGCTAGCGACCTTGTGCCGCCCAGCTGTCCATGGCCCGTGCAGA 2099
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Qy      2100 GCAGAGGACAGTGAGTGTCTGCACTGAGAACCTTAAACCACAGTTGAACATACCCACACC 2159
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Qy      2160 TGTTTGTCTTAAGCTATAGTGTAAAAACAAAGTTTGGGCTCTGAAAATTTAACTGAAAAA 2219
Db      2521 TGTTTGTCTTAAGCTATAGTGTAAAAACAAAGTTTGGGCTCTGAAAATTTAACTGAAAAA 2580
Qy      2220 GATTTCCTTGTTTTTGTAATAGGTGAGATAAAGTACTTAGATTTATAA-GGCAGCTTCCC 2278
Db      2581 GATTTCCTTGTTTTTGTAATAGGTGAGATAAAGTACTTAGATTTATAAGGGCAGCTTCCC 2640
Qy      2279 CTGTAGTGATAAATTACAAGCAGACAATCTTATTTTGTAATGTGATGAAGTGATGATGTC 2338
Db      2641 CTGTAGTGATAAATTACAAGCAGACAATCTTATTTTGTAATGTGATGAAGTGATGATGTC 2700
Qy      2339 TTAACTCTACTTAGAGAGTGTATGTCTGTCTAACAGAACAAAAAGATGCTCTGTGTAAAT 2398
Db      2701 TTAACTCTACTTAGAGAGTGTATGTCTGTCTAACAGAACAAAAAGATGCTCTGTGTAAAT 2760
Qy      2399 TCCTTCCTGTAGGGCACACTGCAGGATTTCCATGTAGATAGAAGAACTATAGGGCCTAGT 2458
Db      2761 TCCTTCCTGTAGGGCACACTGCAGGATTTCCATGTAGATAGAAGAACTATAGGGCCTAGT 2820
Qy      2459 ACAGAAGGTGCACA-CAAATGTTGGCAAAGTC-AAACCCCATGAATTAAAACCTACTGGA 2516
Db      2821 ACAGAAGGTGCACAGCAAATGTTGGCAAAGTCAAAACCCCATGAATTAAAACCTACTGGA 2880
Qy      2517 ATTTGGTTTTTAGGAGTTTGGTAATTAGATTATCTCTTTTGTTATTTTCATTCAGTTATA 2576
Db      2881 ATTTGGTTTTTAGGAGTTTGGTAATTAGATTATCTCTTTTGTTATTTTCATTCAGTTATA 2940
Qy      2577 TCCTTT-GGCTCAGCTAGCTTTGAAATTGGCTGATGAAAAAATATACATAAAAGGGTAAA 2635
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Qy      2636 ATTCACACATACAGCAAACAAAAATGCACAAAGCCTGCTTCGTAACTTTTTTTTCTGGAA 2695
Db      3001 ATTCACACATACAGCAAACAAAAATGCACAAAGCCTGCTTCGTAACTTTTTTTTCTGGAA 3060
Qy      2696 TTGTTTTTCACTTTGCCTTTTTCTGCCAAAACAATAATCAAAGAACTCTTGCTTTAACCT 2755
Db      3061 TTGTTTTTCACTTTGCCTTTTTCTGCCAAAACAATAATCAAAGAACTCTTGCTTTAACCT 3120
Qy      2756 ATTCCTGTACAAAGACTGTTTTTGACCAGATAATCATCTGTTGTGGCATTCTATCTTGTA 2815
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Qy      2876 GTGCCCTGGGAGTCTTAAAAGCAGTGCTTAGCAACATTGGTGATAGCATGTGGCTGGGAC 2935
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Qy      2936 CCAGGGCCCTTCCCCACTCTTCAGCCCCGAGTCATGTGTCTGAGGTGACGGACTGAGACG 2995
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Db      3421 CTGTTTCTTCAAGTACACACTGACTCTGCTACTTTAGGATAAATATATTTTACTCAGAAC 3480
Qy      3116 TCTGAATTTCACAGTATACTTACTAAACTAAGTAAAAATGATACTTAAAATACTTATTTT 3175
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Db      3541 ACTTTCTAGACCTAGGCTAGATGTTTTAAGCTACAGCTCTAGTTCATTGTGATATTTATA 3600
Qy      3236 ATTTGAAAGCTATGAGAATAGATGTGTGGGTGAAGCCATAGAACATATTTGCTTGAAATT 3295
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Qy      3296 CTTGAGCAGGGATCTTATAAAGGGCCAGAAATAAGATGTGTGGTTCACATAGATAGTGAG 3355
Db      3661 CTTGAGCAGGGATCTTATAAAGGGCCAGAAATAAGATGTGTGGTTCACATAGATAGTGAG 3720
Qy      3356 CGTAACATCTGTATTAAACATAGGAGAGAAGTTTATAAAGGGCATTGGCAATAAACTCTT 3415
Db      3721 CGTAACATCTGTATTAAACATAGGAGAGAAGTTTATAAAGGGCATTGGCAATAAACTCTT 3780
Qy      3416 TGTTGCAGCTGTTTTCCAAGCAGTGTAAATACTTTTTCCTGTGATTATGTATAGCCTTGG 3475
Db      3781 TGTTGCAGCTGTTTTCCAAGCAGTGTAAATACTTTTTCCTGTGATTATGTATAGCCTTGG 3840
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Db      3841 AATGGCACCTTTTAACTAACCCATATGTGTTTGGTTTCAATGGTTTTTTATATTCAGATG 3900
Qy      3536 TATATATGGTGCTCACTTTAGGATCAGCAGTGTTGACCATTTATGCTGCATAGCTGTATT 3595
Db      3901 TATATATGGTGCTCACTTTAGGATCAGCAGTGTTGACCATTTATGCTGCATAGCTGTATT 3960
Qy      3596 ATAGCCTTATTAGTTGTGTGGTTGACCCTTGGGGTATACAAATGTCAGTCTGAGTGGTGT 3655
Db      3961 ATAGCCTTATTAGTTGTGTGGTTGACCCTTGGGGTATACAAATGTCAGTCTGAGTGGTGT 4020
Qy      3656 CTTACTCCTTTGTTTATAAGTGAATGATTGTGCATGTTTTGTATGTCATAGTATGTCGTC 3715
Db      4021 CTTACTCCTTTGTTTATAAGTGAATGATTGTGCATGTTTTGTATGTCATAGTATGTCGTC 4080
Qy      3716 ACATAAAAGGGAGGGAGCGAAAAACCATTACATTAAGATAATATTGGACCAAACTACTTA 3775
Db      4081 ACATAAAAGGGAGGGAGCGAAAAACCATTACATTAAGATAATATTGGACCAAACTACTTA 4140
Qy      3776 CTTGCTCTAAACAGTTACTTGTACCCCTTAACCTGTCTTCAAAAGTTGCATATAGTTACA 3835
Db      4141 CTTGCTCTAAACAGTTACTTGTACCCCTTAACCTGTCTTCAAAAGTTGCATATAGTTACA 4200
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Qy       3836 GTAGTGTATAAATTAAATATTGTGGAAAAACAGTCTTGTATTTTTCTGTATGTGTG--TA 3893
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Db       4201 GTAGTGTATAAATTAAATATTGTGGAAAAACAGTCTTGTATTTTTCTGTATGTGTGTATA 4260

Qy       3894 TATATATATAATTATGTACTTCTGGCAATTCTATCTGTATTTAAAGATGTGACAATCTTG 3953
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Db       4261 TATATATATAATTATGTACTTCTGGCAATTCTATCTGTATTTAAAGATGTGACAATCTTG 4320

Qy       3954 ACACCAATTTTAAGAATAGCTGTGAGACCGAATTAAAGATAATCCCTACCAAGTGAAAAT 4013
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Db       4321 ACACCAATTTTAAGAATAGCTGTGAGACCGAATTAAAGATAATCCCTACCAAGTGAAAAT 4380

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Db       4381 TGATGTGTGTTAAGAGGGTACAGAATTATCAACTGATTTGGTCAGTTGCTTCCAATGCTG 4440

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RESULT 7
US-10-311-034-47
; Sequence 47, Application US/10311034
; Publication No. US20040023242A1
; GENERAL INFORMATION:
;  APPLICANT: INCYTE GENOMICS, INC.
;  APPLICANT:  YUE, Henry
;  APPLICANT:  LAL, Preeti
;  APPLICANT:  BANDMAN, Olga
;  APPLICANT:  BOROWSKY, Mark L.
;  APPLICANT:  AU-YOUNG, Janice
;  APPLICANT:  LU, Yan
;  APPLICANT:  GANDHI, Ameena R.
;  APPLICANT:  TRIBOULEY, Catherine M.
;  APPLICANT:  CHAWLA, Narinder K.
;  APPLICANT:  YAO, Monique G.
;  APPLICANT:  LU, Dyung Aina M.
;  APPLICANT:  GREENWALD, Sara R.
;  APPLICANT:  RAMKUMAR, Jayalaxmi
;  APPLICANT:  GRIFFIN, Jennifer A.
;  APPLICANT:  KEARNEY, Liam
;  APPLICANT:  BURFORD, Neil
;  APPLICANT:  NGUYEN, Danniel B.
;  APPLICANT:  TANG, Y. Tom
;  APPLICANT:  BAUGHN, Mariah R.
;  APPLICANT:  HE, Ann
;  APPLICANT:  THORNTON, Michael
;  APPLICANT:  HAFALIA, April
;  APPLICANT:  ARVIZU, Chandra S.
;  APPLICANT:  GURURAJAN, Rajagopal
;  APPLICANT:  LO, Terence P.
;  APPLICANT:  KHAH, Farrah A.
;  APPLICANT:  RECIPON, Shirley A.
;  APPLICANT:  AZIMZAI, Yalda
;  APPLICANT:  POLICKY, Jennifer L.
;  APPLICANT:  DING, Li
;  APPLICANT:  GRETHER, Megan
;  APPLICANT:  ELLIOTT, Vicki S.
;  APPLICANT:  THANGAVELU, Kavitha
;  APPLICANT:  BATRA, Sajeev
;  APPLICANT:  ISON, Craig H.
;  TITLE OF INVENTION: HUMAN KINASES
;  FILE REFERENCE: PI-0125 PCT
;  CURRENT APPLICATION NUMBER: US/10/311,034
;  CURRENT FILING DATE:  2002-12-10
;  PRIOR APPLICATION NUMBER: 60/212,073; 60/213,467; 60/215,651; 60/216,605; 60/218,372;
;    60/228,056
;  PRIOR FILING DATE: 2000-06-15; 2000-06-23; 2000-06-30; 2000-07-07; 2000-07-13; 2000-
;    25
;  NUMBER OF SEQ ID NOS: 52
;  SOFTWARE: PERL Program
; SEQ ID NO 47
;    LENGTH: 4772
;    TYPE: DNA
;    ORGANISM: Homo sapiens
;    FEATURE:
;    NAME/KEY: misc_feature
;    OTHER INFORMATION: Incyte ID No. US20040023242A1 063497CB1
US-10-311-034-47

  Query Match             87.0%;  Score 3622.6;  DB 17;  Length 4772;
  Best Local Similarity   99.7%;  Pred. No. 0;
  Matches 3639;  Conservative    0;  Mismatches    9;  Indels    1;  Gaps    1;

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Qy        241 AACCAACAGGTATAACCATATCACAGCCACATACTTCCTGCTGGCTGAAAGGATCCTGAG 300
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Qy        301 AGAAAAGCAAGAGAAAGAAATACAGACCAGATCTGCAAGCCCGAGCAATATCAAGGCCCA 360
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Db       1630 ATCCAGGAAGAGTGCGCCCGTCCTCAACCAGATCTTTGAGGAAGGGGAATCTGACGATGA 1689

Qy        781 GTTTGACATGGATGAGAATCTGCCTCCCAAGTTGAGCAGGTTAAAGATGAATATAGCTTC 840
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
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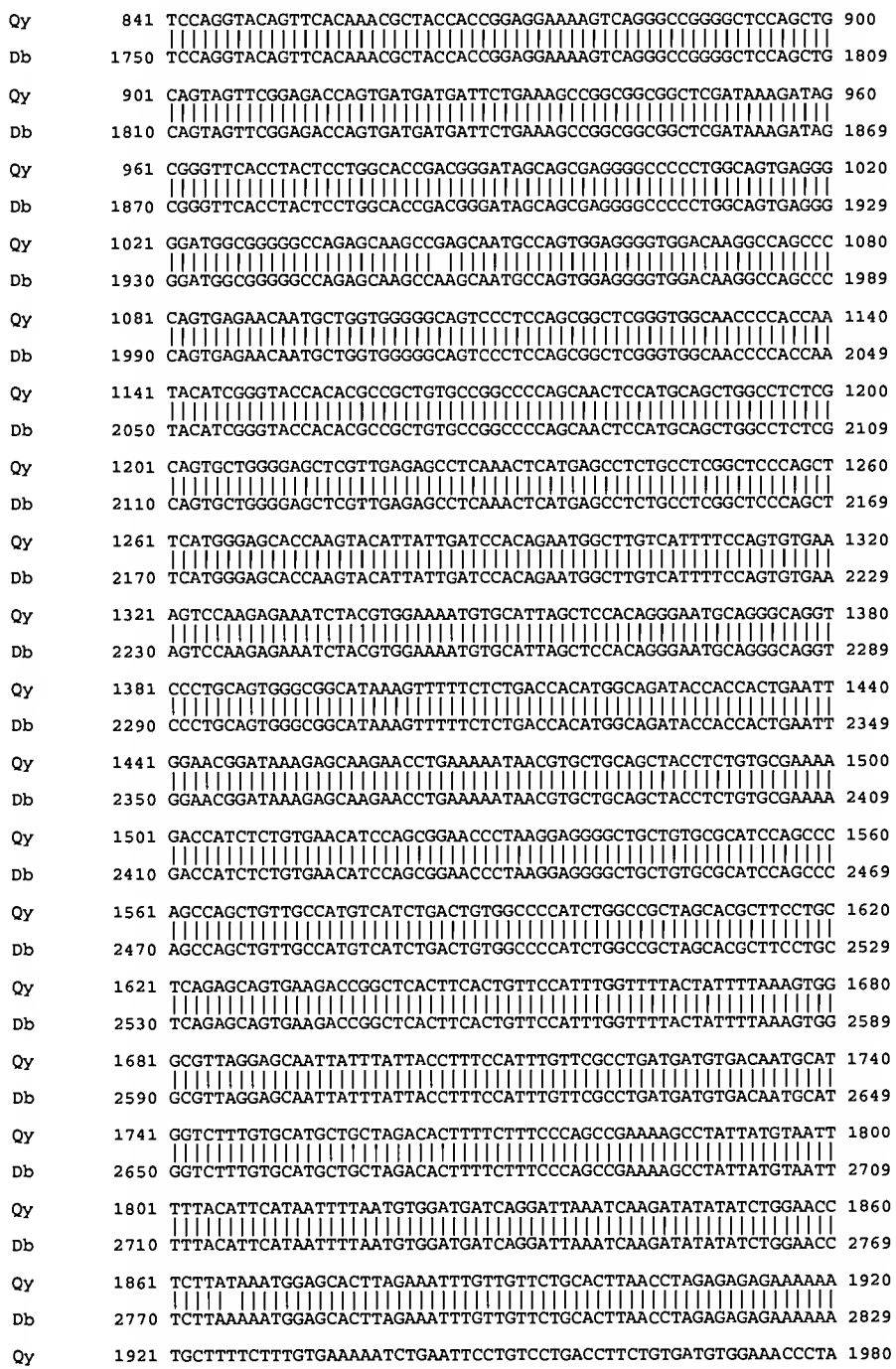
Qy 841 TCCAGGTACAGTTCACAAACGCTACCACCGGAGGAAAAGTCAGGGCCGGGGCTCCAGCTG 900
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Db 2050 TACATCGGGTACCACACGCCGCTGTGCCGGCCCCAGCAACTCCATGCAGCTGGCCTCTCG 2109
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Qy 1261 TCATGGGAGCACCAAGTACATTATTGATCCACAGAATGGCTTGTCATTTTCCAGTGTGAA 1320
Db 2170 TCATGGGAGCACCAAGTACATTATTGATCCACAGAATGGCTTGTCATTTTCCAGTGTGAA 2229
Qy 1321 AGTCCAAGAGAAATCTACGTGGAAAATGTGCATTAGCTCCACAGGGAATGCAGGGCAGGT 1380
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Qy 1921 TGCTTTTCTTTGTGAAAAATCTGAATTCCTGTCCTGACCTTCTGTGATGTGGAAACCCTA 1980
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Qy 1981 GGCTCTGAGACACACTCTCTGGTGTCTGAGACAGAACCAAAGCAATAACGTTGTGATGCC 2040
Db 2890 GGCTCTGAGACACACTCTCTGGTGTCTGAGACAGAACCAAAGCAATAACGTTGTGATGCC 2949
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Qy 2161 GTTTGTCTTAAGCTATAGTGTAAAAACAAAGTTTGGGCTCTGAAAATTTAACTGAAAAAG 2220
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Qy 2221 ATTTCCTTGTTTTTGTAATAGGTGAGATAAAGTACTTAGATTTATAAGGCAGCTTCCCCT 2280
Db 3130 ATTTCCTTGTTTTTGTAATAGGTGAGATAAAGTACTTAGATTTATAAGGCAGCTTCCCCT 3189
Qy 2281 GTAGTGATAAATTACAAGCAGACAATCTTATTTTGTAATGTGATGAAGTGATGATGTCTT 2340
Db 3190 GTAGTGATAAATTACAAGCAGACAATCTTATTTTGTAATGTGATGAAGTGATGATGTCTT 3249
Qy 2341 AACTCTACTTAGAGAGTGTATGTCTGTCTAACAGAACAAAAAGATGCTCTGTGTAAATTC 2400
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Qy 2401 CTTCCTGTAGGGCACACTGCAGGATTTCCATGTAGATAGAAGAACTATAGGGCCTAGTAC 2460
Db 3310 CTTCCTGTAGGGCACACTGCAGGATTTCCATGTAGATAGAAGAACTATAGGGCCTAGTAC 3369
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Db 3370 AGAAGGTGCACACAAATGTTGGCAAAGTCAAAACCCCATGAATTAAAACCTACTGGAATT 3429
Qy 2520 TGGTTTTTAGGAGTTTGGTAATTAGATTATCTCTTTTGTTATTTTCATTCAGTTATATCC 2579
Db 3430 TGGTTTTTAGGAGTTTGGTAATTAGATTATCTCTTTTGTTATTTTCATTCAGTTATATCC 3489
Qy 2580 TTTGGCTCAGCTAGCTTTGAAATTGGCTGATGAAAAAATATACATAAAAGGGTAAAATTC 2639
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Qy 2760 CTGTACAAAGACTGTTTTTGACCAGATAATCATCTGTTGTGGCATTCTATCTTGTAGGAC 2819
Db 3670 CTGTACAAAGACTGTTTTTGACCAGATAATCATCTGTTGTGGCATTCTATCTTGTAGGAC 3729
Qy 2820 ACTGTATATTGCAAATTGCTGATTATGGAAGGGGCCAGTTGCTGTTTTTTCATGCAGTGC 2879
Db 3730 ACTGTATATTGCAAATTGCTGATTATGGAAGGGGCCAGTTGCTGTTTTTTCATGCAGTGC 3789
Qy 2880 CCTGGGAGTCTTAAAAGCAGTGCTTAGCAACATTGGTGATAGCATGTGGCTGGGACCCAG 2939
Db 3790 CCTGGGAGTCTTAAAAGCAGTGCTTAGCAACATTGGTGATAGCATGTGGCTGGGACCCAG 3849
Qy 2940 GGCCCTTCCCCACTCTTCAGCCCCGAGTCATGTGTCTGAGGTGACGGACTGAGACGCATC 2999
Db 3850 GGCCCTTCCCCACTCTTCAGCCCCGAGTCATGTGTCTGAGGTGACGGACTGAGACGCATC 3909
Qy 3000 TGGTCCTGTAATTCAGAGAGTGGGCACATCACCAAAGAACTGCATTGCTGTGGTCACTGT 3059

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Db       3910 TGGTCCTGTAATTCAGAGAGTGGGCACATCACCAAAGAACTGCATTGCTGTGGTCACTGT 3969

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Qy       3120 AATTTCACAGTATACTTACTAAACTAAGTAAAAATGATACTTAAAATACTTATTTTACTT 3179
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       4030 AATTTCACAGTATACTTACTAAACTAAGTAAAAATGATACTTAAAATACTTATTTTACTT 4089

Qy       3180 TCTAGACCTAGGCTAGATGTTTTAAGCTACAGCTCTAGTTCATTGTGATATTTATAATTT 3239
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Db       4090 TCTAGACCTAGGCTAGATGTTTTAAGCTACAGCTCTAGTTCATTGTGATATTTATAATTT 4149

Qy       3240 GAAAGCTATGAGAATAGATGTGTGGGTGAAGCCATAGAACATATTTGCTTGAAATTCTTG 3299
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Db       4150 GAAAGCTATGAGAATAGATGTGTGGGTGAAGCCATAGAACATATTTGCTTGAAATTCTTG 4209

Qy       3300 AGCAGGGATCTTATAAAGGGCCAGAAATAAGATGTGTGGTTCACATAGATAGTGAGCGTA 3359
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Db       4210 AGCAGGGATCTTATAAAGGGCCAGAAATAAGATGTGTGGTTCACATAGATAGTGAGCGTA 4269

Qy       3360 ACATCTGTATTAAACATAGGAGAGAAGTTTATAAAGGGCATTGGCAATAAACTCTTTGTT 3419
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Db       4390 GCACCTTTTAACTAACCCATATGTGTTTGGTTTCAATGGTTTTTTATATTCAGATGTATA 4449

Qy       3540 TATGGTGCTCACTTTAGGATCAGCAGTGTTGACCATTTATGCTGCATAGCTGTATTATAG 3599
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RESULT 8
US-09-925-302-242
; Sequence 242, Application US/09925302
; Patent No. US20020044941A1
; GENERAL INFORMATION:
; APPLICANT: Rosen et al.
; TITLE OF INVENTION: Nucleic Acids, Proteins and Antibodies
; FILE REFERENCE: PA104
; CURRENT APPLICATION NUMBER: US/09/925,302
; CURRENT FILING DATE: 2001-08-10
; PRIOR APPLICATION NUMBER: PCT/US00/05918
; PRIOR FILING DATE: 2000-03-08
; PRIOR APPLICATION NUMBER: 60/124,270
; PRIOR FILING DATE: 1999-03-12
; NUMBER OF SEQ ID NOS: 896
; SOFTWARE: PatentIn Ver. 2.0
; SEQ ID NO 242
; LENGTH: 3276
; TYPE: DNA
; ORGANISM: Homo sapiens
; FEATURE:
; NAME/KEY: misc feature
; LOCATION: (125)
; OTHER INFORMATION: n equals a,t,g, or c
; NAME/KEY: misc feature
; LOCATION: (455)
; OTHER INFORMATION: n equals a,t,g, or c
; NAME/KEY: misc feature
; LOCATION: (1014)
; OTHER INFORMATION: n equals a,t,g, or c
; NAME/KEY: misc feature
; LOCATION: (3276)
; OTHER INFORMATION: n equals a,t,g, or c
US-09-925-302-242

Query Match 76.2%; Score 3174.6; DB 9; Length 3276;
Best Local Similarity 99.4%; Pred. No. 0;
Matches 3237; Conservative 8; Mismatches 4; Indels 6; Gaps 6;

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Db          2 GACTAGTGATGATGATTCTGAAAGCCGGCGGCGGCTCGATAAAGATAGCGGGTTCACCTA 61

Qy        973 CTCCTGGCACCGACGGGATAGCAGCGAGGGGCCCCCTGGCAGTGAGGGGGATGGCGGGGG 1032
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db         62 CTCCTGGCACCGACGGGATAGCAGCGAGGGGCCCCCTGGCAGTGAGGGGGATGGCGGGGG 121

Qy       1033 CCAGAGCAAGCCGAGCAATGCCAGTGGAGGGGTGGACAAGGCCAGCCCCAGTGAGAACAA 1092
              ||| |||||||| ||||||||||||||||||||||||||||||||||||||||||||||||
Db         122 CCANAGCAAGCCAAGCAATGCCAGTGGAGGGGTGGACAAGGCCAGCCCCAGTGAGAACAA 181

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Db         301 GCTCGTTGAGAGCCTCAAACTCATGAGCCTCTGCCTCGGCTCCCAGCTTCATGGGAGCAC 360

Qy       1273 CAAGTACATTATTGATCCACAGAATGGCTTGTCATTTTCCAGTGTGAAAGTCCAAGAGAA 1332
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              |||||||||||||||||||||||||| ||||||||||||||||||||||||||||||||
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Db         719 AGACCGGCTCACTTCACTGTTCCATTTGGTTTTACTATTTTAAAGTGGGCGTTAGGAGCA 778

Qy       1693 ATTATTTATTACCTTTCCATTTGTTCGCCTGATGATGTGACAATGCATGGTCTTTGTGCA 1752
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Db         839 TGCTGCTAGACACTTTWMTTTCCCAGCCGAAAAGCCTATTATGTAATTTTTACATTCATA 898

Qy       1813 ATTTTAATGTGGATGATCAGGATTAAATCAAGATATATATCTGGAACCTCTTATAAATGG 1872
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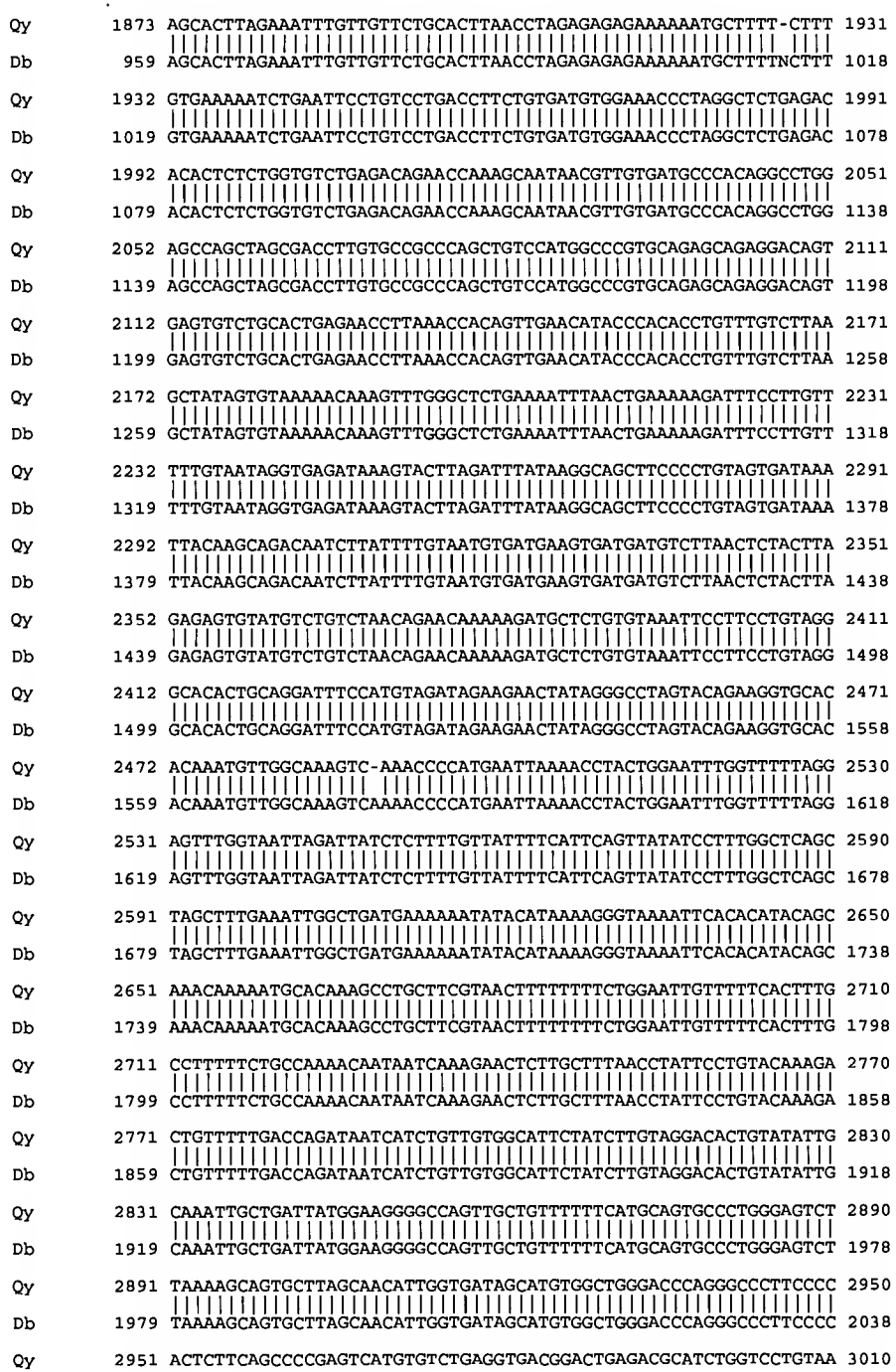
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Qy 1992 ACACTCTCTGGTGTCTGAGACAGAACCAAAGCAATAACGTTGTGATGCCCACAGGCCTGG 2051
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Qy 2052 AGCCAGCTAGCGACCTTGTGCCGCCCAGCTGTCCATGGCCCGTGCAGAGCAGAGGACAGT 2111
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Qy 2112 GAGTGTCTGCACTGAGAACCTTAAACCACAGTTGAACATACCCACACCTGTTTGTCTTAA 2171
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Qy 2172 GCTATAGTGTAAAAACAAAGTTTGGGCTCTGAAAATTTAACTGAAAAAGATTTCCTTGTT 2231
Db 1259 GCTATAGTGTAAAAACAAAGTTTGGGCTCTGAAAATTTAACTGAAAAAGATTTCCTTGTT 1318
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Db 1379 TTACAAGCAGACAATCTTATTTTGTAATGTGATGAAGTGATGATGTCTTAACTCTACTTA 1438
Qy 2352 GAGAGTGTATGTCTGTCTAACAGAACAAAAAGATGCTCTGTGTAAATTCCTTCCTGTAGG 2411
Db 1439 GAGAGTGTATGTCTGTCTAACAGAACAAAAAGATGCTCTGTGTAAATTCCTTCCTGTAGG 1498
Qy 2412 GCACACTGCAGGATTTCCATGTAGATAGAAGAACTATAGGGCCTAGTACAGAAGGTGCAC 2471
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Db 1559 ACAAATGTTGGCAAAGTCAAAACCCCATGAATTAAAACCTACTGGAATTTGGTTTTTAGG 1618
Qy 2531 AGTTTGGTAATTAGATTATCTCTTTTGTTATTTTCATTCAGTTATATCCTTTGGCTCAGC 2590
Db 1619 AGTTTGGTAATTAGATTATCTCTTTTGTTATTTTCATTCAGTTATATCCTTTGGCTCAGC 1678
Qy 2591 TAGCTTTGAAATTGGCTGATGAAAAAATATACATAAAAGGGTAAAATTCACACATACAGC 2650
Db 1679 TAGCTTTGAAATTGGCTGATGAAAAAATATACATAAAAGGGTAAAATTCACACATACAGC 1738
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Db 2099 TTCAGAGAGTGGGCACATCACCAAAGAACTGCATTGCTGTGGTCACTGTTTCTTCAAGTA 2158
Qy 3071 CACACTGACTCTGCTACTTTAGGATAAATATATTTTACTCAGAACTCTGAATTTCACAGT 3130
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Db 2279 GCTAGATGTTTTAAGCTACAGCTCTAGTTCATTGTGATATTTATAATTTGAAAGCTATGA 2338
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Qy 3311 TATAAAGGGCCAGAAATAAGATGTGTGGTTCACATAGATAGTGAGCGTAACATCTGTATT 3370
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Db 2459 AAACATAGGAGAGAAGTTTATAAAGGGCATTGGCAATAAACTCTTTGTTGCAGCTGTTTT 2518
Qy 3431 CCAAGCAGTGTAAATACTTTTTCCTGTGATTATGTATAGCCTTGGAATGGCACCTTTTAA 3490
Db 2519 CCAAGCAGTGTAAATACTTTTTCCTGTGATTATGTATAGCCTTGGAATGGCACCTTTTAA 2578
Qy 3491 CTAACCCATATGTGTTTGGTTTCAATGGTTTTTTATATTCAGATGTATATATGGTGCTCA 3550
Db 2579 CTAACCCATATGTGTTTGGTTTCAATGGTTTTTTATATTCAGATGTATATAKGGTGCTCA 2638
Qy 3551 CTTTAGGATCAGCAGTGTTGACCATTTATGCTGCATAGCTGTATTATAGCCTTATTAGTT 3610
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Qy 3611 GTGTGGTTGACCCTTGGGGTATACAAATGTCAGTCTGAGTGGTGTCTTACTCCTTTGTTT 3670
Db 2699 GTGTGGTTGACCCTTGGGGTATACAAATGTCAGTCTGAGTGGTGTCTTACTCC-TTGKTT 2757
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Db 2998 ACTTCTGGCAATTCTATCTGTATTTAAAGATGTGACAATCTTGACACCAATTTTAAGAAT 3057
Qy 3971 AGCTGTGAGACCGAATTAAAGATAATCCCTACCAAGTGAAAATTGATGTGTGTTAAGAGG 4030
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Db       3118 GTACAGAATTATCAACTGATTTGGTCAGTTGCTTCCAATGCTGGTTGATTTCCCTCATTG 3177

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RESULT 9
US-09-925-302-242
; Sequence 242, Application US/09925302
; Publication No. US20030064072A9
; GENERAL INFORMATION:
;  APPLICANT: Rosen et al.
;  TITLE OF INVENTION: Nucleic Acids, Proteins and Antibodies
;  FILE REFERENCE: PA104
;  CURRENT APPLICATION NUMBER: US/09/925,302
;  CURRENT FILING DATE:  2001-08-10
;  PRIOR APPLICATION NUMBER: PCT/US00/05918
;  PRIOR FILING DATE: 2000-03-08
;  PRIOR APPLICATION NUMBER: 60/124,270
;  PRIOR FILING DATE: 1999-03-12
;  NUMBER OF SEQ ID NOS: 896
;  SOFTWARE: PatentIn Ver. 2.0
; SEQ ID NO 242
;   LENGTH: 3276
;   TYPE: DNA
;   ORGANISM: Homo sapiens
;   FEATURE:
;   NAME/KEY: misc feature
;   LOCATION: (125)
;   OTHER INFORMATION: n equals a,t,g, or c
;   NAME/KEY: misc feature
;   LOCATION: (455)
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;   LOCATION: (1014)
;   OTHER INFORMATION: n equals a,t,g, or c
;   NAME/KEY: misc feature
;   LOCATION: (3276)
;   OTHER INFORMATION: n equals a,t,g, or c
US-09-925-302-242

  Query Match             76.2%;  Score 3174.6;  DB 10;  Length 3276;
  Best Local Similarity   99.4%;  Pred. No. 0;
  Matches 3237;  Conservative    8;  Mismatches    4;  Indels    6;  Gaps    6;

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Db        421 ATYTACGTGGAAAATGTGCATTAGCTCCACAGGGNATGCAGGGCAGGTCCCT-CAGTGGG 479

Qy       1393 CGGCATAAAGTTTTTCTCTGACCACATGGCAGATACCACCACTGAATTGGAACGGATAAA 1452
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        480 CGGCATAAAGTTTTTCTCTGACCACATGGCAGATACCACCACTGAATTGGAACGGATAAA 539

Qy       1453 GAGCAAGAACCTGAAAAATAACGTGCTGCAGCTACCTCTGTGCGAAAAGACCATCTCTGT 1512
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        540 GAGCAAGAACCTGAAAAATAACGTGCTGCAGCTACCTCTGTGCGAAAAGACCATCTCTGT 599

Qy       1513 GAACATCCAGCGGAACCCTAAGGAGGGGCTGCTGTGCGCATCCAGCCCAGCCAGCTGTTG 1572
              |||||||||||||||||||||||||||||| ||||||||||||||||||||||||||||
Db        600 GAACATCCAGCGGAACCCTAAGGAGGGGCT-CTGTGCGCATCCAGCCCAGCCAGCTGTTG 658

Qy       1573 CCATGTCATCTGACTGTGGCCCCATCTGGCCGCTAGCACGCTTCCTGCTCAGAGCAGTGA 1632
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        659 CCATGTCATCTGACTGTGGCCCCATCTGGCCGCTAGCACGCTTCCTGCTCAGAGCAGTGA 718

Qy       1633 AGACCGGCTCACTTCACTGTTCCATTTGGTTTTACTATTTTAAAGTGGGCGTTAGGAGCA 1692
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        719 AGACCGGCTCACTTCACTGTTCCATTTGGTTTTACTATTTTAAAGTGGGCGTTAGGAGCA 778

Qy       1693 ATTATTTATTACCTTTCCATTTGTTCGCCTGATGATGTGACAATGCATGGTCTTTGTGCA 1752
              |||||||||||||||||||||||:||||||||||||||||||||||||||||||||||||
Db        779 ATTATTTATTACCTTTCCATTTGTWCGCCTGATGATGTGACAATGCATGGTCTTTGTGCA 838

Qy       1753 TGCTGCTAGACACTTTTCTTTCCCAGCCGAAAAGCCTATTATGTAATTTTTACATTCATA 1812
              ||||||||||||||::||||||||||||||||||||||||||||||||||||||||||||
Db        839 TGCTGCTAGACACTTTWMTTTCCCAGCCGAAAAGCCTATTATGTAATTTTTACATTCATA 898

Qy       1813 ATTTTAATGTGGATGATCAGGATTAAATCAAGATATATATCTGGAACCTCTTATAAATGG 1872
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        899 ATTTTAATGTGGATGATCAGGATTAAATCAAGATATATATCTGGAACCTCTTATAAATGG 958

Qy       1873 AGCACTTAGAAATTTGTTGTTCTGCACTTAACCTAGAGAGAGAAAAAATGCTTTT-CTTT 1931
              |||||||||||||||||||||||||||||||||||||||||||||||||||| |||| ||||
Db        959 AGCACTTAGAAATTTGTTGTTCTGCACTTAACCTAGAGAGAGAAAAAATGCTTTTNCTTT 1018

Qy       1932 GTGAAAAATCTGAATTCCTGTCCTGACCTTCTGTGATGTGGAAACCCTAGGCTCTGAGAC 1991
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        1019 GTGAAAAATCTGAATTCCTGTCCTGACCTTCTGTGATGTGGAAACCCTAGGCTCTGAGAC 1078

Qy       1992 ACACTCTCTGGTGTCTGAGACAGAACCAAAGCAATAACGTTGTGATGCCCACAGGCCTGG 2051
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        1079 ACACTCTCTGGTGTCTGAGACAGAACCAAAGCAATAACGTTGTGATGCCCACAGGCCTGG 1138

Qy       2052 AGCCAGCTAGCGACCTTGTGCCGCCCAGCTGTCCATGGCCCGTGCAGAGCAGAGGACAGT 2111
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        1139 AGCCAGCTAGCGACCTTGTGCCGCCCAGCTGTCCATGGCCCGTGCAGAGCAGAGGACAGT 1198

Qy       2112 GAGTGTCTGCACTGAGAACCTTAAACCACAGTTGAACATACCCACACCTGTTTGTCTTAA 2171
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        1199 GAGTGTCTGCACTGAGAACCTTAAACCACAGTTGAACATACCCACACCTGTTTGTCTTAA 1258

Qy       2172 GCTATAGTGTAAAAACAAAGTTTGGGCTCTGAAAATTTAACTGAAAAAGATTTCCTTGTT 2231
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        1259 GCTATAGTGTAAAAACAAAGTTTGGGCTCTGAAAATTTAACTGAAAAAGATTTCCTTGTT 1318

Qy       2232 TTTGTAATAGGTGAGATAAAGTACTTAGATTTATAAGGCAGCTTCCCCTGTAGTGATAAA 2291
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        1319 TTTGTAATAGGTGAGATAAAGTACTTAGATTTATAAGGCAGCTTCCCCTGTAGTGATAAA 1378

Qy       2292 TTACAAGCAGACAATCTTATTTTGTAATGTGATGAAGTGATGATGTCTTAACTCTACTTA 2351
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        1379 TTACAAGCAGACAATCTTATTTTGTAATGTGATGAAGTGATGATGTCTTAACTCTACTTA 1438
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Qy       2352 GAGAGTGTATGTCTGTCTAACAGAACAAAAAGATGCTCTGTGTAAATTCCTTCCTGTAGG 2411
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       1439 GAGAGTGTATGTCTGTCTAACAGAACAAAAAGATGCTCTGTGTAAATTCCTTCCTGTAGG 1498

Qy       2412 GCACACTGCAGGATTTCCATGTAGATAGAAGAACTATAGGGCCTAGTACAGAAGGTGCAC 2471
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       1499 GCACACTGCAGGATTTCCATGTAGATAGAAGAACTATAGGGCCTAGTACAGAAGGTGCAC 1558

Qy       2472 ACAAATGTTGGCAAAGTC-AAACCCCCATGAATTAAAACCTACTGGAATTTGGTTTTTAGG 2530
             |||||||||||||||||| ||||||||||||||||||||||||||||||||||||||||||
Db       1559 ACAAATGTTGGCAAAGTCAAAACCCCATGAATTAAAACCTACTGGAATTTGGTTTTTAGG 1618

Qy       2531 AGTTTGGTAATTAGATTATCTCTTTTGTTATTTTCATTCAGTTATATCCTTTGGCTCAGC 2590
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       1619 AGTTTGGTAATTAGATTATCTCTTTTGTTATTTTCATTCAGTTATATCCTTTGGCTCAGC 1678

Qy       2591 TAGCTTTGAAATTGGCTGATGAAAAAATATACATAAAAGGGTAAAATTCACACATACAGC 2650
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       1679 TAGCTTTGAAATTGGCTGATGAAAAAATATACATAAAAGGGTAAAATTCACACATACAGC 1738

Qy       2651 AAACAAAAATGCACAAAGCCTGCTTCGTAACTTTTTTTTCTGGAATTGTTTTTCACTTTG 2710
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       1739 AAACAAAAATGCACAAAGCCTGCTTCGTAACTTTTTTTTCTGGAATTGTTTTTCACTTTG 1798

Qy       2711 CCTTTTTCTGCCAAAACAATAATCAAAGAACTCTTGCTTTAACCTATTCCTGTACAAAGA 2770
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       1799 CCTTTTTCTGCCAAAACAATAATCAAAGAACTCTTGCTTTAACCTATTCCTGTACAAAGA 1858

Qy       2771 CTGTTTTTGACCAGATAATCATCTGTTGTGGCATTCTATCTTGTAGGACACTGTATATTG 2830
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       1859 CTGTTTTTGACCAGATAATCATCTGTTGTGGCATTCTATCTTGTAGGACACTGTATATTG 1918

Qy       2831 CAAATTGCTGATTATGGAAGGGGCCAGTTGCTGTTTTTTCATGCAGTGCCCTGGGAGTCT 2890
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       1919 CAAATTGCTGATTATGGAAGGGGCCAGTTGCTGTTTTTTCATGCAGTGCCCTGGGAGTCT 1978

Qy       2891 TAAAAGCAGTGCTTAGCAACATTGGTGATAGCATGTGGCTGGGACCCAGGGCCCTTCCCC 2950
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       1979 TAAAAGCAGTGCTTAGCAACATTGGTGATAGCATGTGGCTGGGACCCAGGGCCCTTCCCC 2038

Qy       2951 ACTCTTCAGCCCCGAGTCATGTGTCTGAGGTGACGGACTGAGACGCATCTGGTCCTGTAA 3010
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       2039 ACTCTTCAGCCCCGAGTCATGTGTCTGAGGTGACGGACTGAGACGCATCTGGTCCTGTAA 2098

Qy       3011 TTCAGAGAGTGGGCACATCACCAAAGAACTGCATTGCTGTGGTCACTGTTTCTTCAAGTA 3070
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       2099 TTCAGAGAGTGGGCACATCACCAAAGAACTGCATTGCTGTGGTCACTGTTTCTTCAAGTA 2158

Qy       3071 CACACTGACTCTGCTACTTTAGGATAAATATATTTTACTCAGAACTCTGAATTTCACAGT 3130
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       2159 CACACTGACTCTGCTACTTTAGGATAAATATATTTTACTCAGAACTCTGAATTTCACAGT 2218

Qy       3131 ATACTTACTAAACTAAGTAAAAATGATACTTAAAATACTTATTTTACTTTCTAGACCTAG 3190
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       2219 ATACTTACTAAACTAAGTAAAAATGATACTTAAAATACTTATTTTACTTTCTAGACCTAG 2278

Qy       3191 GCTAGATGTTTTAAGCTACAGCTCTAGTTCATTGTGATATTTATAATTTGAAAGCTATGA 3250
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       2279 GCTAGATGTTTTAAGCTACAGCTCTAGTTCATTGTGATATTTATAATTTGAAAGCTATGA 2338

Qy       3251 GAATAGATGTGTGGGTGAAGCCATAGAACATATTTGCTTGAAATTCTTGAGCAGGGATCT 3310
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       2339 GAATAGATGTGTGGGTGAAGCCATAGAACATATTTGCTTGAAATTCTTGAGCAGGGATCT 2398

Qy       3311 TATAAAGGGCCAGAAATAAGATGTGTGGTTCACATAGATAGTGAGCGTAACATCTGTATT 3370
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       2399 TATAAAGGGCCAGAAATAAGATGTGTGGTTCACATAGATAGTGAGCGTAACATCTGTATT 2458

Qy       3371 AAACATAGGAGAGAAGTTTATAAAGGGCATTGGCAATAAACTCTTTGTTGCAGCTGTTTT 3430
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       2459 AAACATAGGAGAGAAGTTTATAAAGGGCATTGGCAATAAACTCTTTGTTGCAGCTGTTTT 2518

Qy       3431 CCAAGCAGTGTAAATACTTTTTCCTGTGATTATGTATAGCCTTGGAATGGCACCTTTTAA 3490
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       2519 CCAAGCAGTGTAAATACTTTTTCCTGTGATTATGTATAGCCTTGGAATGGCACCTTTTAA 2578

Qy       3491 CTAACCCATATGTGTTTGGTTTCAATGGTTTTTTATATTCAGATGTATATATGGTGCTCA 3550
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||:|||||||
Db       2579 CTAACCCATATGTGTTTGGTTTCAATGGTTTTTTATATTCAGATGTATATAKGGTGCTCA 2638

Qy       3551 CTTTAGGATCAGCAGTGTTGACCATTTATGCTGCATAGCTGTATTATAGCCTTATTAGTT 3610
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       2639 CTTTAGGATCAGCAGTGTTGACCATTTATGCTGCATAGCTGTATTATAGCCTTATTAGTT 2698

Qy       3611 GTGTGGTTGACCCTTGGGGTATACAAATGTCAGTCTGAGTGGTGTCTTACTCCTTTGTTT 3670
             |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||| |||:||
Db       2699 GTGTGGTTGACCCTTGGGGTATACAAATGTCAGTCTGAGTGGTGTCTTACTCC-TTGKTT 2757

Qy       3671 ATAAGTGAATGATTGTGCATGTTTTGTATGTCATAGTATGTCGTCACATAAAAGGGAGGG 3730
             |||||||||||||||||||||||||||:|||||:||||||||||||||||||||||||||||||
Db       2758 ATAAGTGAATGATTGTGCATGTTTKGTATGYCATAGTATGTCGTCACATAAAAGGGAGGG 2817

Qy       3731 AGCGAAAAACCATTACATTAAGATAATATTGGACCAAACTACTTACTTGCTCTAAACAGT 3790
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       2818 AGCGAAAAACCATTACATTAAGATAATATTGGACCAAACTACTTACTTGCTCTAAACAGT 2877

Qy       3791 TACTTGTACCCCTTAACCTGTCTTCAAAAGTTGCATATAGTTACAGTAGTGTATAAATTA 3850
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       2878 TACTTGTACCCCTTAACCTGTCTTCAAAAGTTGCATATAGTTACAGTAGTGTATAAATTA 2937

Qy       3851 AATATTGTGGAAAAAACAGTCTTGTATTTTTCTGTATGTGTGTATATATATATAATTATGT 3910
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       2938 AATATTGTGGAAAAAACAGTCTTGTATTTTTCTGTATGTGTGTATATATATATAATTATGT 2997

Qy       3911 ACTTCTGGCAATTCTATCTGTATTTAAAGATGTGACAATCTTGACACCAATTTTAAGAAT 3970
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       2998 ACTTCTGGCAATTCTATCTGTATTTAAAGATGTGACAATCTTGACACCAATTTTAAGAAT 3057

Qy       3971 AGCTGTGAGACCGAATTAAAGATAATCCCTACCAAGTGAAAATTGATGTGTGTTAAGAGG 4030
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       3058 AGCTGTGAGACCGAATTAAAGATAATCCCTACCAAGTGAAAATTGATGTGTGTTAAGAGG 3117

Qy       4031 GTACAGAATTATCAACTGATTTGGTCAGTTGCTTCCAATGCTGGTTGATTTCCCTCATTG 4090
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       3118 GTACAGAATTATCAACTGATTTGGTCAGTTGCTTCCAATGCTGGTTGATTTCCCTCATTG 3177

Qy       4091 TGTAAACATTGACAGGTATGTGACAAATGGGAAAAAAAATCCAAATAATAAAGTGACATA 4150
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       3178 TGTAAACATTGACAGGTATGTGACAAATGGGAAAAAAAATCCAAATAATAAAGTGACATA 3237

Qy       4151 TTGGTGTTCAGCAAT 4165
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Db       3238 TTGGTGTTCAGCAAT 3252
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RESULT 10
US-10-108-260A-2237
; Sequence 2237, Application US/10108260A
; Publication No. US20040005560A1
; GENERAL INFORMATION:
; APPLICANT: HELIX RESEARCH INSTITUTE
; TITLE OF INVENTION: No. US20040005560A1el full length cDNA
; FILE REFERENCE: H1-A0106
; CURRENT APPLICATION NUMBER: US/10/108,260A
; CURRENT FILING DATE: 2002-03-27
; NUMBER OF SEQ ID NOS: 5458
; SOFTWARE: PatentIn Ver. 2.1
; SEQ ID NO 2237
; LENGTH: 1768
; TYPE: DNA
; ORGANISM: Homo sapiens
US-10-108-260A-2237

Query Match 23.1%; Score 962.4; DB 17; Length 1768;
Best Local Similarity 99.4%; Pred. No. 7.7e-243;
Matches 966; Conservative 0; Mismatches 6; Indels 0; Gaps 0;

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Qy       364 TAGGCAGTCATGGCCAACCAAAATTGATGTACCCCAGGACCTTGAGGATGACCTCACGGC 423
            ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       556 TAGGCAGTCATGGCCAACCAAAATTGATGTACCCCAGGACCTTGAGGATGACCTCACGGC 615

Qy       424 CACTCCTTTGTCCCACGCGACTGTCCCTCAGTCTCCTGCTCGGGCTGCTGACAGTGTCCT 483
            ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       616 CACTCCTTTGTCCCACGCGACTGTCCCTCAGTCTCCTGCTCGGGCTGCTGACAGTGTCCT 675

Qy       484 CAATGGCCACAGGAGCAAAGGCCTGTGTGACTCAGCTAAGAAAGATGACCTCCCTGAGTT 543
            ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       676 CAATGGCCACAGGAGCAAAGGCCTGTGTGACTCAGCTAAGAAAGATGACCTCCCTGAGTT 735

Qy       544 GGCTGGACCAGCACTCTCTACGGTGCCACCCGCAAGCTTAAAACCCACAGCCAGTGGGCG 603
            ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       736 GGCTGGACCAGCACTCTCTACGGTGCCACCCGCAAGCTTAAAACCCACAGCCAGTGGGCG 795

Qy       604 GAAGTGTCTGTTCAGGGTGGAAGAAGATGAAGAGGAAGATGAGGAGGACAAGAAACCCAT 663
            ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       796 GAAGTGTCTGTTCAGGGTGGAAGAAGATGAAGAGGAAGATGAGGAGGACAAGAAACCCAT 855

Qy       664 GTCCCTCTCAACACAAGTGGTTTTGCGCCGGAAGCCATCTGTAACCAACCGCCTGACATC 723
            ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       856 GTCCCTCTCAACACAAGTGGTTTTGCGCCGGAAGCCATCTGTAACCAACCGCCTGACATC 915

Qy       724 CAGGAAGAGTGCGCCCGTCCTCAACCAGATCTTTGAGGAAGGGGAATCTGATGATGAGTT 783
            ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       916 CAGGAAGAGTGCGCCCGTCCTCAACCAGATCTTTGAGGAAGGGGAATCTGACGATGAGTT 975

Qy       784 TGACATGGATGAGAATCTGCCTCCCAAGTTGAGCAGGTTAAAGATGAATATAGCTTCTCC 843
            ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       976 TGACATGGATGAGAATCTGCCTCCCAAGTTGAGCAGGTTAAAGATGAATATAGCTTCTCC 1035

Qy       844 AGGTACAGTTCACAAACGCTACCACCGGAGGAAAAGTCAGGGCCGGGGCTCCAGCTGCAG 903
            ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db      1036 AGGTACAGTTCACAAACGCTACCACCGGAGGAAAAGTCAGGGCCGGGGCTCCAGCTGCAG 1095

Qy       904 TAGTTCGGAGACCAGTGATGATGATTCTGAAAGCCGGCGGCGGCTCGATAAAGATAGCGG 963
            ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db      1096 TAGTTCGGAGACCAGTGATGATGATTCTGAAAGCCGGCGGCGGCTCGATAAAGATAGCGG 1155

Qy       964 GTTCACCTACTCCTGGCACCGACGGGATAGCAGCGAGGGGCCCCCTGGCAGTGAGGGGGA 1023
            ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db      1156 GTTCACCTACTCCTGGCACCGACGGGATAGCAGCGAGGGGCCCCCTGGCAGTGAGGGGGA 1215

Qy      1024 TGGCGGGGGCCAGAGCAAGCCGAGCAATGCCAGTGGAGGGGTGGACAAGGCCAGCCCCAG 1083
            ||||||||||||||||||||| ||||||||||||||||||||||||||||||||||||||||
Db      1216 TGGCGGGGGCCAGAGCAAGCCAAGCAATGCCAGTGGAGGGGTGGACAAGGCCAGCCCCAG 1275

Qy      1084 TGAGAACAATGCTGGTGGGGGCAGTCCCTCCAGCGGCTCGGGTGGCAACCCCACCAATAC 1143
            ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db      1276 TGAGAACAATGCTGGTGGGGGCAGTCCCTCCAGCGGCTCGGGTGGCAACCCCACCAATAC 1335

Qy      1144 ATCGGGTACCACACGCCGCTGTGCCGGCCCCAGCAACTCCATGCAGCTGGCCTCTCGCAG 1203
            ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db      1336 ATCGGGTACCACACGCCGCTGTGCCGGCCCCAGCAACTCCATGCAGCTGGCCTCTCGCAG 1395

Qy      1204 TGCTGGGGAGCTCGTTGAGAGCCTCAAACTCATGAGCCTCTGCCTCGGCTCCCAGCTTCA 1263
            ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db      1396 TGCTGGGGAGCTCGTTGAGAGCCTCAAACTCATGAGCCTCTGCCTCGGCTCCCAGCTTCA 1455

Qy      1264 TGGGAGCACCAAGTACATTATTGATCCACAGAATGGCTTGTCATTTTCCAGTGTGAAAGT 1323
            ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db      1456 TGGGAGCACCAAGTACATTATTGATCCACAGAATGGCTTGTCATTTTCCAGTGTGAAAGT 1515

Qy      1324 CCAAGAGAAATC 1335
            ||||   | |||
Db      1516 CCAAATTACATC 1527
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RESULT 11
US-09-918-995-23795
; Sequence 23795, Application US/09918995
; Publication No. US20030073623A1
; GENERAL INFORMATION:
; APPLICANT: Hyseq, Inc.
; TITLE OF INVENTION: NOVEL NUCLEIC ACID SEQUENCES OBTAINED
; TITLE OF INVENTION: FROM VARIOUS cDNA LIBRARIES
; FILE REFERENCE: 20411-756
; CURRENT APPLICATION NUMBER: US/09/918,995
; CURRENT FILING DATE: 2001-07-30
; PRIOR APPLICATION NUMBER: US/09/235,076
; PRIOR FILING DATE: 1999-01-20
; NUMBER OF SEQ ID NOS: 38054
; SOFTWARE: FastSEQ for Windows Version 3.0
; SEQ ID NO 23795
; LENGTH: 502
; TYPE: DNA
; ORGANISM: Homo sapiens
; FEATURE:
; NAME/KEY: misc_feature
; LOCATION: (1)...(502)
; OTHER INFORMATION: n = A,T,C or G
US-09-918-995-23795

Query Match 10.9%; Score 454.4; DB 10; Length 502;
Best Local Similarity 97.5%; Pred. No. 6.3e-109;
Matches 461; Conservative 0; Mismatches 12; Indels 0; Gaps 0;

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Qy      2888 TCTTAAAAGCAGTGCTTAGCAACATTGGTGATAGCATGTGGCTGGGACCCAGGGCCCTTC 2947
            ||  | ||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        29 TCGCACGAGCAGTGCTTAGCAACATTGGTGATAGCATGTGGCTGGGACCCAGGGCCCTTC 88

Qy      2948 CCCACTCTTCAGCCCCGAGTCATGTGTCTGAGGTGACGGACTGAGACGCATCTGGTCCTG 3007
            ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        89 CCCACTCTTCAGCCCCGAGTCATGTGTCTGAGGTGACGGACTGAGACGCATCTGGTCCTG 148

Qy      3008 TAATTCAGAGAGTGGGCACATCACCAAAGAACTGCATTGCTGTGGTCACTGTTTCTTCAA 3067
            ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       149 TAATTCAGAGAGTGGGCACATCACCAAAGAACTGCATTGCTGTGGTCACTGTTTCTTCAA 208

Qy      3068 GTACACACTGACTCTGCTACTTTAGGATAAATATATTTTACTCAGAACTCTGAATTTCAC 3127
            ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       209 GTACACACTGACTCTGCTACTTTAGGATAAATATATTTTACTCAGAACTCTGAATTTCAC 268

Qy      3128 AGTATACTTACTAAACTAAGTAAAAATGATACTTAAAATACTTATTTTACTTTCTAGACC 3187
            ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       269 AGTATACTTACTAAACTAAGTAAAAATGATACTTAAAATACTTATTTTACTTTCTAGACC 328

Qy      3188 TAGGCTAGATGTTTTAAGCTACAGCTCTAGTTCATTGTGATATTTATAATTTGAAAGCTA 3247
            ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       329 TAGGCTAGATGTTTTAAGCTACAGCTCTAGTTCATTGTGATATTTATAATTTGAAAGCTA 388

Qy      3248 TGAGAATAGATGTGTGGGTGAAGCCATAGAACATATTTGCTTGAAATTCTTGAGCAGGGA 3307
            ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       389 TGAGAATAGATGTGTGGGTGAAGCCATAGAACATATTTGCTTGAAATTCTTGAGCAGGGA 448

Qy      3308 TCTTATAAAGGGCCAGAAATAAGATGTGTGGTTCACATAGATAGTGAGCGTAA 3360
            ||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       449 TCTTATAAAGGGCCAGAAATAAGATGTGTGGTTCACATNAGATAGTAGCGTAA 501
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RESULT 12
US-09-969-034-4185/c
; Sequence 4185, Application US/09969034
; Publication No. US20040110668A1
; GENERAL INFORMATION:
; APPLICANT: Burgess, Christopher C.
; APPLICANT: Astle, Jon H.
; APPLICANT: Carroll, Eddie III
; APPLICANT: Catino, Theodore J.
; APPLICANT: Dwivedi, Poornima
; APPLICANT: Molino, Gary A.
; APPLICANT: Thiagalingam, Arunthathi
; APPLICANT: Lewis, Marcia E.
; TITLE OF INVENTION: Nucleic Acid Sequences Differentially

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;  TITLE OF INVENTION:  Expressed in Cancer Tissue
;  FILE REFERENCE: 1657/1032
;  CURRENT APPLICATION NUMBER: US/09/969,034
;  CURRENT FILING DATE:  2001-10-02
;  PRIOR APPLICATION NUMBER: 60/237,271
;  PRIOR FILING DATE: 2000-02-10
;  NUMBER OF SEQ ID NOS: 4494
;  SOFTWARE: FastSEQ for Windows Version 4.0
; SEQ ID NO 4185
;   LENGTH: 505
;   TYPE: DNA
;   ORGANISM: Homo sapiens
;   FEATURE:
;   NAME/KEY: misc_feature
;   LOCATION: 482, 485, 503
;   OTHER INFORMATION: n = A,T,C or G
US-09-969-034-4185

  Query Match             9.8%;  Score 409.6;  DB 11;  Length 505;
  Best Local Similarity   97.0%;  Pred. No. 4.4e-97;
  Matches  460;  Conservative    0;  Mismatches    9;  Indels    5;  Gaps    4;

Qy       3324 AAATAAGATGTGTGGTTCACATAGATAGTGAGCGTAACATCTGTATTAAACATAGGAGAG 3383
              | |  |||||||||||||||||||| |||||||||||||||||| ||||||||||||
Db        469 AGAATAGATGTGTGGTTCACATAGATA-TGAGCGTAACATCTGTATT-AACATAGGAGAG 412

Qy       3384 AAGTTTATAAAGGGCATTGGCAATAAACTCTTTGTTGCAGCTGTTTTCCAAGCAGTGTAA 3443
              |||||||| ||||||||||||||||||     ||  |||||||||||||||||||||||||
Db        411 AAGTTTAT-AAGGGCATTGGCAATAACTC--TTTGTGCAGCTGTTTTCCAAGCAGTGTAA 355

Qy       3444 ATACTTTTTCCTGTGATTATGTATAGCCTTGGAATGGCACCTTTTAACTAACCCATATGT 3503
              |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        354 ATACTTTTTCCTGTGATTATGTATAGCCTTGGAATGGCACCTTTTAACTAACCCATATGT 295

Qy       3504 GTTTGGTTTCAATGGTTTTTTATATTCAGATGTATATATGGTGCTCACTTTAGGATCAGC 3563
              |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        294 GTTTGGTTTCAATGGTTTTTTATATTCAGATGTATATATGGTGCTCACTTTAGGATCAGC 235

Qy       3564 AGTGTTGACCATTTATGCTGCATAGCTGTATTATAGCCTTATTAGTTGTGTGGTTGACCC 3623
              |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        234 AGTGTTGACCATTTATGCTGCATAGCTGTATTATAGCCTTATTAGTTGTGTGGTTGACCC 175

Qy       3624 TTGGGGTATACAAATGTCAGTCTGAGTGGTGTCTTACTCCTTTGTTTATAAGTGAATGAT 3683
              |||||||||||||||||||||||||||||||||||| |||||||||||||||||||||||
Db        174 TTGGGGTATACAAATGTCAGTCTGAGTGGTGTCTTACTTCTTTGTTTATAAGTGAATGAT 115

Qy       3684 TGTGCATGTTTTGTATGTCATAGTATGTCGTCACATAAAAGGGAGGGAGCGAAAAACCAT 3743
              |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        114 TGTGCATGTTTTGTATGTCATAGTATGTCGTCACATAAAAGGGAGGGAGCGAAAAACCAT 55

Qy       3744 TACATTAAGATAATATTGGACCAAACTACTTACTTGCTCTAAACAGTTACTTGT 3797
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db          54 TACATTAAGATAATATTGGACCAAACTACTTACTTGCTCTAAACAGTTACTTGT 1


RESULT 13
US-09-867-701-3627
; Sequence 3627, Application US/09867701
; Patent No. US20020132237A1
; GENERAL INFORMATION:
;  APPLICANT: Aglate, Paul A.
;  APPLICANT:  Jones, Robert
;  APPLICANT:  Harlocker, Susan L.
;  TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY
;  TITLE OF INVENTION:  AND DIAGNOSIS OF OVARIAN CANCER
;  FILE REFERENCE: 210121.497
;  CURRENT APPLICATION NUMBER: US/09/867,701
;  CURRENT FILING DATE:  2001-05-29
;  NUMBER OF SEQ ID NOS: 10912
;  SOFTWARE: FastSEQ for Windows Version 4.0
; SEQ ID NO 3627
;   LENGTH: 369
;   TYPE: DNA
;   ORGANISM: Homo sapien
US-09-867-701-3627

  Query Match             8.9%;  Score 369;  DB 9;  Length 369;
  Best Local Similarity   100.0%;  Pred. No. 1.9e-86;
  Matches  369;  Conservative    0;  Mismatches    0;  Indels    0;  Gaps    0;

Qy       3111 AGAACTCTGAATTTCACAGTATACTTACTAAACTAAGTAAAAATGATACTTAAAATACTT 3170
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db          1 AGAACTCTGAATTTCACAGTATACTTACTAAACTAAGTAAAAATGATACTTAAAATACTT 60

Qy       3171 ATTTTACTTTCTAGACCTAGGCTAGATGTTTTAAGCTACAGCTCTAGTTCATTGTGATAT 3230
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db         61 ATTTTACTTTCTAGACCTAGGCTAGATGTTTTAAGCTACAGCTCTAGTTCATTGTGATAT 120

Qy       3231 TTATAATTTGAAAGCTATGAGAATAGATGTGTGGGTGAAGCCATAGAACATATTTGCTTG 3290
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        121 TTATAATTTGAAAGCTATGAGAATAGATGTGTGGGTGAAGCCATAGAACATATTTGCTTG 180

Qy       3291 AAATTCTTGAGCAGGGATCTTATAAAGGGCCAGAAATAAGATGTGTGGTTCACATAGATA 3350
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        181 AAATTCTTGAGCAGGGATCTTATAAAGGGCCAGAAATAAGATGTGTGGTTCACATAGATA 240

Qy       3351 GTGAGCGTAACATCTGTATTAAACATAGGAGAGAAGTTTATAAAGGGCATTGGCAATAAA 3410
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        241 GTGAGCGTAACATCTGTATTAAACATAGGAGAGAAGTTTATAAAGGGCATTGGCAATAAA 300

Qy       3411 CTCTTTGTTGCAGCTGTTTTCCAAGCAGTGTAAATACTTTTTCCTGTGATTATGTATAGC 3470
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        301 CTCTTTGTTGCAGCTGTTTTCCAAGCAGTGTAAATACTTTTTCCTGTGATTATGTATAGC 360

Qy       3471 CTTGGAATG 3479
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Db        361 CTTGGAATG 369


RESULT 14
US-09-969-347-122/c
; Sequence 122, Application US/09969347
; Patent No. US20020115085A1
; GENERAL INFORMATION:
;  APPLICANT: Ebner, Reinhard
;  TITLE OF INVENTION: Cancer Gene Determination and Therapeutic Screening Using Signat
;  TITLE OF INVENTION:  Sets
;  FILE REFERENCE: 689290-69
;  CURRENT APPLICATION NUMBER: US/09/969,347
;  CURRENT FILING DATE:  2001-10-02
;  PRIOR APPLICATION NUMBER: US/60/237,598
;  PRIOR FILING DATE: 2000-10-03
;  PRIOR APPLICATION NUMBER: US/60/237,604
;  PRIOR FILING DATE: 2000-10-03
;  NUMBER OF SEQ ID NOS: 318
;  SOFTWARE: PatentIn version 3.0
; SEQ ID NO 122
;   LENGTH: 396
;   TYPE: DNA
;   ORGANISM: Homo sapiens
US-09-969-347-122

  Query Match             7.7%;  Score 320;  DB 9;  Length 396;
  Best Local Similarity   90.5%;  Pred. No. 1.8e-73;
  Matches  362;  Conservative   21;  Mismatches   13;  Indels    4;  Gaps    4;

Qy       3766 AAACTACTTACTTGCTCTAAACAGTTACTTGTACCCCTTAACCTGTCTTCAAAAGTTGCA 3825
              |||||||||:|||||:|||||:|| |||||||:|||||||| ||||||||||||| :|||
Db         396 AAACTACTTWCTTGCYCTAAAMAG-TACTTGTDCCCCTTAACCTGTCTTCAAAAGGKGCA 338

Qy       3826 TATAGTTACAGTAGTGTATAAATTAAATATTGTGGAAAAACAGTCTTGTATTTTTCTGTA 3885
              |||||||||||||||||||||||||||||||||||  |||||||||||||:|||
Db         337 TATAGTTACAGTAGTGTATAAATTAAATATTGTGG-GAAACAGTCTTGTAKTTTCTGTAT 279

Qy       3886 TGTGTGTATATATATATAATTATGTACTTCTGGCAATTCTATCTGTATTTAAAGATGTGA 3945
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Db        278 GTGGSVBAGABATATATAAWTATGTACYTCTGGCAADTCTAWCTGTAWTTAAAGATGTGR 219

Qy       3946 CAATCTTGACACCAATTTTAAGAATAGCTGTGAGACCGAATTAAAGATAATCCCTACCAA 4005
              ||||||||||||||||||||||||||||||||||||||||:||| ||||||||||||||||||
Db        2I8 CAATCTTGACACCAATTTTAAGAATAGCTGTGAGACCGRATT-AAGATAATCCCTACCAA I60

Qy       4006 GTGAAAATTGATGTGTGTTAAGAGGGTACAGAATTATCAACTGATTTGGTCAGTTGCTTC 4065
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        I59 GTGAAAATTGATGTGTGTTAAGAGGGTACAGAATTATCAACTGATTTGGTCAGTTGCTTC 100

Qy       4066 CAATGCTGGTTGATTTCCCTCATTGTGTAAACATTGACAGGTATGTGACAAATGGGAAAA 4I25
              ||||||||||||||||||:|||||||| ||||||||||||||||||||||||||||::
Db         99 CAATGCTGGTTGATTTCCCYCATTGTGT-AACATTGACAGGTATGTGACAAATGGGAARV 4I

Qy       4I26 AAAATCCAAATAATAAAGTGACATATTGGTGTTCAGCAAT 4I65
              :|||:|||||||||||||||||||||||||||||||||||
Db         40 RAAABCCAAATAATAAAGTGACATATTGGTGTTCAGCAAT I


RESULT I5
US-09-880-107-I703/c
; Sequence I703, Application US/09880107
; Patent No. US20020I4298IAI
; GENERAL INFORMATION:
;  APPLICANT: Horne, Darci T.
;  APPLICANT:  Vockley, Joseph G.
;  APPLICANT:  Scherf, Uwe
;  APPLICANT:  Gene Logic, Inc.
;  TITLE OF INVENTION: Gene Expression Profiles in Liver Cancer
;  FILE REFERENCE: 44921-5028-WO
;  CURRENT APPLICATION NUMBER: US/09/880,I07
;  CURRENT FILING DATE:  200I-06-I4
;  PRIOR APPLICATION NUMBER: US 60/2I1,379
;  PRIOR FILING DATE: 2000-06-14
;  PRIOR APPLICATION NUMBER: US 60/237,054
;  PRIOR FILING DATE: 2000-I0-02
;  NUMBER OF SEQ ID NOS: 3950
;  SOFTWARE: PatentIn Ver. 2.I
; SEQ ID NO I703
;   LENGTH: 396
;   TYPE: DNA
;   ORGANISM: Homo sapiens
;   FEATURE:
;   OTHER INFORMATION: Genbank Accession No. US20020I4298IAI D60769
US-09-880-107-1703

  Query Match             7.7%;  Score 320;  DB 9;  Length 396;
  Best Local Similarity   90.5%;  Pred. No. I.8e-73;
  Matches  362;  Conservative   21;  Mismatches   I3;  Indels    4;  Gaps    4;

Qy       3766 AAACTACTTACTTGCTCTAAACAGTTACTTGTACCCCTTAACCTGTCTTCAAAAGTTGCA 3825
              |||||||||:|||||:|||||:|| |||||||:|||||||||||||||||||| :|||
Db         396 AAACTACTTWCTTGCYCTAAAMAG-TACTTGTDCCCCTTAACCTGTCTTCAAAAGGKGCA 338

Qy       3826 TATAGTTACAGTAGTGTATAAATTAAATATTGTGGAAAAACAGTCTTGTATTTTTCTGTA 3885
              |||||||||||||||||||||||||||||||||  |||||||||||||:|||
Db         337 TATAGTTACAGTAGTGTATAAATTAAATATTGTGG-GAAACAGTCTTGTAKTTTCTGTAT 279

Qy       3886 TGTGTGTATATATATATAATTATGTACTTCTGGCAATTCTATCTGTATTTAAAGATGTGA 3945
                 | ::| |:||||||||:|||||||:||||||||:||||:|||||:||||||||||:
Db         278 GTGGSVBAGABATATATAAWTATGTACYTCTGGCAADTCTAWCTGTAWTTAAAGATGTGR 219

Qy       3946 CAATCTTGACACCAATTTTAAGAATAGCTGTGAGACCGAATTAAAGATAATCCCTACCAA 4005
              ||||||||||||||||||||||||||||||||||||||||:||| ||||||||||||||||||
Db         218 CAATCTTGACACCAATTTTAAGAATAGCTGTGAGACCGRATT-AAGATAATCCCTACCAA I60

Qy       4006 GTGAAAATTGATGTGTGTTAAGAGGGTACAGAATTATCAACTGATTTGGTCAGTTGCTTC 4065
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db         I59 GTGAAAATTGATGTGTGTTAAGAGGGTACAGAATTATCAACTGATTTGGTCAGTTGCTTC I00

Qy       4066 CAATGCTGGTTGATTTCCCTCATTGTGTAAACATTGACAGGTATGTGACAAATGGGAAAA 4I25
              ||||||||||||||||||:|||||||| ||||||||||||||||||||||||||||::
Db          99 CAATGCTGGTTGATTTCCCYCATTGTGT-AACATTGACAGGTATGTGACAAATGGGAARV 4I

Qy       4I26 AAAATCCAAATAATAAAGTGACATATTGGTGTTCAGCAAT 4I65
              :|||:|||||||||||||||||||||||||||||||||||
Db          40 RAAABCCAAATAATAAAGTGACATATTGGTGTTCAGCAAT I


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Db        143 ATATATTTATATATTTTTTA 124


RESULT 13
US-09-949-016-54024/c
; Sequence 54024, Application US/09949016
; Patent No. 6812339
; GENERAL INFORMATION:
;  APPLICANT: VENTER, J. Craig et al.
;  TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES ASSOCIATED
;  TITLE OF INVENTION:  WITH HUMAN DISEASE, METHODS OF DETECTION AND USES THEREOF
;  FILE REFERENCE: CL001307
;  CURRENT APPLICATION NUMBER: US/09/949,016
;  CURRENT FILING DATE:  2000-04-14
;  PRIOR APPLICATION NUMBER: 60/241,755
;  PRIOR FILING DATE: 2000-10-20
;  PRIOR APPLICATION NUMBER: 60/237,768
;  PRIOR FILING DATE: 2000-10-03
;  PRIOR APPLICATION NUMBER: 60/231,498
;  PRIOR FILING DATE: 2000-09-08
;  NUMBER OF SEQ ID NOS: 207012
;  SOFTWARE: FastSEQ for Windows Version 4.0
; SEQ ID NO 54024
;   LENGTH: 601
;   TYPE: DNA
;   ORGANISM: Human
US-09-949-016-54024

  Query Match             1.1%;  Score 44;  DB 4;  Length 601;
  Best Local Similarity   57.1%;  Pred. No. 0.072;
  Matches   80;  Conservative    0;  Mismatches   60;  Indels    0;  Gaps    0;

Qy       3826 TATAGTTACAGTAGTGTATAAATTAAATATTGTGGAAAAACAGTCTTGTATTTTTCTGTA 3885
              |||| ||| |    | | || ||  | | || |  | |    |    | ||||||| | ||
Db        202 TATATTTATATATATATTTATATATATTTTTATATATATTTATATGTATATTTTTATATA 143

Qy       3886 TGTGTGTATATATATATAATTATGTACTTCTGGCAATTCTATCTGTATTTAAAGATGTGA 3945
              | | | ||| |||||||| || | ||  | |    |  | ||| | |||||| |  | |
Db        142 TTTATATATTTATATATATTTTTATATATTTATAATATTTATATATATTTATATTTATTT 83

Qy       3946 CAATCTTGACACCAATTTTA 3965
              || || | |    |||||
Db         82 ATATATTTATATATTTTTA 63


RESULT 14
US-09-949-016-108654/c
; Sequence 108654, Application US/09949016
; Patent No. 6812339
; GENERAL INFORMATION:
;  APPLICANT: VENTER, J. Craig et al.
;  TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES ASSOCIATED
;  TITLE OF INVENTION:  WITH HUMAN DISEASE, METHODS OF DETECTION AND USES THEREOF
;  FILE REFERENCE: CL001307
;  CURRENT APPLICATION NUMBER: US/09/949,016
;  CURRENT FILING DATE:  2000-04-14
;  PRIOR APPLICATION NUMBER: 60/241,755
;  PRIOR FILING DATE: 2000-10-20
;  PRIOR APPLICATION NUMBER: 60/237,768
;  PRIOR FILING DATE: 2000-10-03
;  PRIOR APPLICATION NUMBER: 60/231,498
;  PRIOR FILING DATE: 2000-09-08
;  NUMBER OF SEQ ID NOS: 207012
;  SOFTWARE: FastSEQ for Windows Version 4.0
; SEQ ID NO 108654
;   LENGTH: 601
;   TYPE: DNA
;   ORGANISM: Human
US-09-949-016-108654

  Query Match             1.1%;  Score 44;  DB 4;  Length 601;
  Best Local Similarity   57.1%;  Pred. No. 0.072;
  Matches   80;  Conservative    0;  Mismatches   60;  Indels    0;  Gaps    0;

Qy       3826 TATAGTTACAGTAGTGTATAAATTAAATATTGTGGAAAAACAGTCTTGTATTTTTCTGTA 3885
              |||| ||| |    | | || ||  | | || |  | |    |    | ||||||| | ||
Db        285 TATATTTATATATATATTTATATATATTTTTATATATATTTATATGTATATTTTTATATA 226

Qy       3886 TGTGTGTATATATATATAATTATGTACTTCTGGCAATTCTATCTGTATTTAAAGATGTGA 3945
              | | | ||| |||||||| || | ||  | |    |  | ||| | |||||| |  | |
Db        225 TTTATATATTTATATATATTTTTATATATTTATAATATTTATATATATTTATATTTATTT 166

Qy       3946 CAATCTTGACACCAATTTTA 3965
              || || | |    |||||
Db        165 ATATATTTATATATTTTTA 146


RESULT 15
US-09-949-016-108655/c
; Sequence 108655, Application US/09949016
; Patent No. 6812339
; GENERAL INFORMATION:
;  APPLICANT: VENTER, J. Craig et al.
;  TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES ASSOCIATED
;  TITLE OF INVENTION:  WITH HUMAN DISEASE, METHODS OF DETECTION AND USES THEREOF
;  FILE REFERENCE: CL001307
;  CURRENT APPLICATION NUMBER: US/09/949,016
;  CURRENT FILING DATE:  2000-04-14
;  PRIOR APPLICATION NUMBER: 60/241,755
;  PRIOR FILING DATE: 2000-10-20
;  PRIOR APPLICATION NUMBER: 60/237,768
;  PRIOR FILING DATE: 2000-10-03
;  PRIOR APPLICATION NUMBER: 60/231,498
;  PRIOR FILING DATE: 2000-09-08
;  NUMBER OF SEQ ID NOS: 207012
;  SOFTWARE: FastSEQ for Windows Version 4.0
; SEQ ID NO 108655
;   LENGTH: 601
;   TYPE: DNA
;   ORGANISM: Human
US-09-949-016-108655

  Query Match             1.1%;  Score 44;  DB 4;  Length 601;
  Best Local Similarity   57.1%;  Pred. No. 0.072;
  Matches   80;  Conservative    0;  Mismatches   60;  Indels    0;  Gaps    0;

Qy       3826 TATAGTTACAGTAGTGTATAAATTAAATATTGTGGAAAAACAGTCTTGTATTTTTCTGTA 3885
              |||| ||| |    | | || ||  | | || |  | |    |    | ||||||| | ||
Db        272 TATATTTATATATATATTTATATATATTTTTATATATATTTATATGTATATTTTTATATA 213

Qy       3886 TGTGTGTATATATATATAATTATGTACTTCTGGCAATTCTATCTGTATTTAAAGATGTGA 3945
              | | | ||| |||||||| || | ||  | |    |  | ||| | |||||| |  | |
Db        212 TTTATATATTTATATATATTTTTATATATTTATAATATTTATATATATTTATATTTATTT 153

Qy       3946 CAATCTTGACACCAATTTTA 3965
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Db        152 ATATATTTATATATTTTTA 133


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Job time : 632.842 secs
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;  SOFTWARE: FastSEQ for Windows Version 4.0
; SEQ ID NO 54020
;   LENGTH: 601
;   TYPE: DNA
;   ORGANISM: Human
US-09-949-016-54020

  Query Match             1.1%;  Score 44;  DB 4;  Length 601;
  Best Local Similarity   57.1%;  Pred. No. 0.072;
  Matches   80;  Conservative    0;  Mismatches   60;  Indels    0;  Gaps    0;

Qy       3826 TATAGTTACAGTAGTGTATAAATTAAATATTGTGGAAAAACAGTCTTGTATTTTTCTGTA 3885
              |||| ||| |    | | || ||  | | || |  | |   |    | |||||||| | ||
Db        285 TATATTTATATATATATTTATATATATTTTTATATATATTTATATGTATATTTTTATATA 226

Qy       3886 TGTGTGTATATATATATAATTATGTACTTCTGGCAATTCTATCTGTATTTAAAGATGTGA 3945
              | | | ||| |||||||| || | ||  | |   | | ||| | |||||| |  | |
Db        225 TTTATATATTTATATATATTTTTATATATTTATAATATTTATATATATTTATATTTATTT 166

Qy       3946 CAATCTTGACACCAATTTTA 3965
                || || | |    |||||
Db        165 ATATATTTATATATTTTTTA 146


RESULT 10
US-09-949-016-54021/c
; Sequence 54021, Application US/09949016
; Patent No. 6812339
; GENERAL INFORMATION:
;  APPLICANT: VENTER, J. Craig et al.
;  TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES ASSOCIATED
;  TITLE OF INVENTION:  WITH HUMAN DISEASE, METHODS OF DETECTION AND USES THEREOF
;  FILE REFERENCE: CL001307
;  CURRENT APPLICATION NUMBER: US/09/949,016
;  CURRENT FILING DATE:  2000-04-14
;  PRIOR APPLICATION NUMBER: 60/241,755
;  PRIOR FILING DATE: 2000-10-20
;  PRIOR APPLICATION NUMBER: 60/237,768
;  PRIOR FILING DATE: 2000-10-03
;  PRIOR APPLICATION NUMBER: 60/231,498
;  PRIOR FILING DATE: 2000-09-08
;  NUMBER OF SEQ ID NOS: 207012
;  SOFTWARE: FastSEQ for Windows Version 4.0
; SEQ ID NO 54021
;   LENGTH: 601
;   TYPE: DNA
;   ORGANISM: Human
US-09-949-016-54021

  Query Match             1.1%;  Score 44;  DB 4;  Length 601;
  Best Local Similarity   57.1%;  Pred. No. 0.072;
  Matches   80;  Conservative    0;  Mismatches   60;  Indels    0;  Gaps    0;

Qy       3826 TATAGTTACAGTAGTGTATAAATTAAATATTGTGGAAAAACAGTCTTGTATTTTTCTGTA 3885
              |||| ||| |    | | || ||  | | || |  | |   |    | |||||||| | ||
Db        272 TATATTTATATATATATTTATATATATTTTTATATATATTTATATGTATATTTTTATATA 213

Qy       3886 TGTGTGTATATATATATAATTATGTACTTCTGGCAATTCTATCTGTATTTAAAGATGTGA 3945
              | | | ||| |||||||| || | ||  | |   | | ||| | |||||| |  | |
Db        212 TTTATATATTTATATATATTTTTATATATTTATAATATTTATATATATTTATATTTATTT 153

Qy       3946 CAATCTTGACACCAATTTTA 3965
                || || | |    |||||
Db        152 ATATATTTATATATTTTTTA 133


RESULT 11
US-09-949-016-54022/c
; Sequence 54022, Application US/09949016
; Patent No. 6812339
; GENERAL INFORMATION:
;  APPLICANT: VENTER, J. Craig et al.
;  TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES ASSOCIATED
;  TITLE OF INVENTION:  WITH HUMAN DISEASE, METHODS OF DETECTION AND USES THEREOF
;  FILE REFERENCE: CL001307
;  CURRENT APPLICATION NUMBER: US/09/949,016
;  CURRENT FILING DATE:  2000-04-14
;  PRIOR APPLICATION NUMBER: 60/241,755
;  PRIOR FILING DATE: 2000-10-20
;  PRIOR APPLICATION NUMBER: 60/237,768
;  PRIOR FILING DATE: 2000-10-03
;  PRIOR APPLICATION NUMBER: 60/231,498
;  PRIOR FILING DATE: 2000-09-08
;  NUMBER OF SEQ ID NOS: 207012
;  SOFTWARE: FastSEQ for Windows Version 4.0
; SEQ ID NO 54022
;   LENGTH: 601
;   TYPE: DNA
;   ORGANISM: Human
US-09-949-016-54022

  Query Match             1.1%;  Score 44;  DB 4;  Length 601;
  Best Local Similarity   57.1%;  Pred. No. 0.072;
  Matches   80;  Conservative    0;  Mismatches   60;  Indels    0;  Gaps    0;

Qy       3826 TATAGTTACAGTAGTGTATAAATTAAATATTGTGGAAAAACAGTCTTGTATTTTTCTGTA 3885
              |||| ||| |    | | || ||  | | || |  | |   |    | |||||||| | ||
Db        267 TATATTTATATATATATTTATATATATTTTTATATATATTTATATGTATATTTTTATATA 208

Qy       3886 TGTGTGTATATATATATAATTATGTACTTCTGGCAATTCTATCTGTATTTAAAGATGTGA 3945
              | | | ||| |||||||| || | ||  | |   | | ||| | |||||| |  | |
Db        207 TTTATATATTTATATATATTTTTATATATTTATAATATTTATATATATTTATATTTATTT 148

Qy       3946 CAATCTTGACACCAATTTTA 3965
                || || | |    |||||
Db        147 ATATATTTATATATTTTTTA 128


RESULT 12
US-09-949-016-54023/c
; Sequence 54023, Application US/09949016
; Patent No. 6812339
; GENERAL INFORMATION:
;  APPLICANT: VENTER, J. Craig et al.
;  TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES ASSOCIATED
;  TITLE OF INVENTION:  WITH HUMAN DISEASE, METHODS OF DETECTION AND USES THEREOF
;  FILE REFERENCE: CL001307
;  CURRENT APPLICATION NUMBER: US/09/949,016
;  CURRENT FILING DATE:  2000-04-14
;  PRIOR APPLICATION NUMBER: 60/241,755
;  PRIOR FILING DATE: 2000-10-20
;  PRIOR APPLICATION NUMBER: 60/237,768
;  PRIOR FILING DATE: 2000-10-03
;  PRIOR APPLICATION NUMBER: 60/231,498
;  PRIOR FILING DATE: 2000-09-08
;  NUMBER OF SEQ ID NOS: 207012
;  SOFTWARE: FastSEQ for Windows Version 4.0
; SEQ ID NO 54023
;   LENGTH: 601
;   TYPE: DNA
;   ORGANISM: Human
US-09-949-016-54023

  Query Match             1.1%;  Score 44;  DB 4;  Length 601;
  Best Local Similarity   57.1%;  Pred. No. 0.072;
  Matches   80;  Conservative    0;  Mismatches   60;  Indels    0;  Gaps    0;

Qy       3826 TATAGTTACAGTAGTGTATAAATTAAATATTGTGGAAAAACAGTCTTGTATTTTTCTGTA 3885
              |||| ||| |    | | || ||  | | || |  | |   |    | |||||||| | ||
Db        263 TATATTTATATATATATTTATATATATTTTTATATATATTTATATGTATATTTTTATATA 204

Qy       3886 TGTGTGTATATATATATAATTATGTACTTCTGGCAATTCTATCTGTATTTAAAGATGTGA 3945
              | | | ||| |||||||| || | ||  | |   | | ||| | |||||| |  | |
Db        203 TTTATATATTTATATATATTTTTATATATTTATAATATTTATATATATTTATATTTATTT 144

Qy       3946 CAATCTTGACACCAATTTTA 3965
```

```
;  TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES ASSOCIATED
;  TITLE OF INVENTION:  WITH HUMAN DISEASE, METHODS OF DETECTION AND USES THEREOF
;  FILE REFERENCE: CL001307
;  CURRENT APPLICATION NUMBER: US/09/949,016
;  CURRENT FILING DATE:  2000-04-14
;  PRIOR APPLICATION NUMBER: 60/241,755
;  PRIOR FILING DATE: 2000-10-20
;  PRIOR APPLICATION NUMBER: 60/237,768
;  PRIOR FILING DATE: 2000-10-03
;  PRIOR APPLICATION NUMBER: 60/231,498
;  PRIOR FILING DATE: 2000-09-08
;  NUMBER OF SEQ ID NOS: 207012
;  SOFTWARE: FastSEQ for Windows Version 4.0
; SEQ ID NO 61868
;   LENGTH: 582
;   TYPE: DNA
;   ORGANISM: Human
US-09-949-016-61868

  Query Match             1.1%;  Score 45.4;  DB 4;  Length 582;
  Best Local Similarity   58.5%;  Pred. No. 0.027;
  Matches   79;  Conservative    0;  Mismatches   56;  Indels    0;  Gaps    0;

Qy       3817 AAAGTTGCATATAGTTACAGTAGTGTATAAATTAAATATTGTGGAAAAACAGTCTTGTAT 3876
             | | ||  ||||| ||| |     | |||| ||  | |  | |  |   | |    | |||
Db        184 ATATTTTTATATATTTATATATATTTATATATATATTTATATATATTTATATATATTTAT 125

Qy       3877 TTTTCTGTATGTGTGTATATATATATAATTATGTACTTCTGGCAATTCTATCTGTATTTA 3936
             | | | ||| | | ||| |||||||| |||| || || |    ||| ||| | ||||||
Db        124 ATATATTTATATATATATTTATATATATTTATATATTTATATATATTTTATATATATTTA 65

Qy       3937 AAGATGTGACAATCT 3951
             | || |    || |
Db          64 TATATATTTATATAT 50


RESULT 7
US-09-949-016-13539/c
; Sequence 13539, Application US/09949016
; Patent No. 6812339
; GENERAL INFORMATION:
;  APPLICANT: VENTER, J. Craig et al.
;  TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES ASSOCIATED
;  TITLE OF INVENTION:  WITH HUMAN DISEASE, METHODS OF DETECTION AND USES THEREOF
;  FILE REFERENCE: CL001307
;  CURRENT APPLICATION NUMBER: US/09/949,016
;  CURRENT FILING DATE:  2000-04-14
;  PRIOR APPLICATION NUMBER: 60/241,755
;  PRIOR FILING DATE: 2000-10-20
;  PRIOR APPLICATION NUMBER: 60/237,768
;  PRIOR FILING DATE: 2000-10-03
;  PRIOR APPLICATION NUMBER: 60/231,498
;  PRIOR FILING DATE: 2000-09-08
;  NUMBER OF SEQ ID NOS: 207012
;  SOFTWARE: FastSEQ for Windows Version 4.0
; SEQ ID NO 13539
;   LENGTH: 314798
;   TYPE: DNA
;   ORGANISM: Human
;   FEATURE:
;   NAME/KEY: misc_feature
;   LOCATION: (1)...(314798)
;   OTHER INFORMATION: n = A,T,C or G
US-09-949-016-13539

  Query Match             1.1%;  Score 45.4;  DB 4;  Length 314798;
  Best Local Similarity   58.5%;  Pred. No. 2.7;
  Matches   79;  Conservative    0;  Mismatches   56;  Indels    0;  Gaps    0;

Qy       3817 AAAGTTGCATATAGTTACAGTAGTGTATAAATTAAATATTGTGGAAAAACAGTCTTGTAT 3876
             | | ||  ||||| ||| |     | |||| ||  | |  | |  |   | |    | |||
Db      19013 ATATTTTTATATATTTATATATATTTATATATATATTTATATATATTTATATATATTTAT 18954

Qy       3877 TTTTCTGTATGTGTGTATATATATATAATTATGTACTTCTGGCAATTCTATCTGTATTTA 3936
             | | | ||| | | ||| |||||||| |||| || || |    ||| ||| | ||||||
Db      18953 ATATATTTATATATATATTTATATATATTTATATATTTATATATATTTTATATATATTTA 18894

Qy       3937 AAGATGTGACAATCT 3951
             | || |    || |
Db      18893 TATATATTTATATAT 18879


RESULT 8
US-09-949-016-13432
; Sequence 13432, Application US/09949016
; Patent No. 6812339
; GENERAL INFORMATION:
;  APPLICANT: VENTER, J. Craig et al.
;  TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES ASSOCIATED
;  TITLE OF INVENTION:  WITH HUMAN DISEASE, METHODS OF DETECTION AND USES THEREOF
;  FILE REFERENCE: CL001307
;  CURRENT APPLICATION NUMBER: US/09/949,016
;  CURRENT FILING DATE:  2000-04-14
;  PRIOR APPLICATION NUMBER: 60/241,755
;  PRIOR FILING DATE: 2000-10-20
;  PRIOR APPLICATION NUMBER: 60/237,768
;  PRIOR FILING DATE: 2000-10-03
;  PRIOR APPLICATION NUMBER: 60/231,498
;  PRIOR FILING DATE: 2000-09-08
;  NUMBER OF SEQ ID NOS: 207012
;  SOFTWARE: FastSEQ for Windows Version 4.0
; SEQ ID NO 13432
;   LENGTH: 76553
;   TYPE: DNA
;   ORGANISM: Human
US-09-949-016-13432

  Query Match             1.1%;  Score 44.8;  DB 4;  Length 76553;
  Best Local Similarity   49.2%;  Pred. No. 1.4;
  Matches  118;  Conservative    0;  Mismatches  122;  Indels    0;  Gaps    0;

Qy       3826 TATAGTTACAGTAGTGTATAAATTAAATATTGTGGAAAAACAGTCTTGTATTTTTCTGTA 3885
             || |||    |   | ||| || |||  | ||  ||  |   | || |||||||| |
Db      53125 TACAGTGGTTGGCATATATTAAGTAATAAATGGTGATTATTGGCTTTTTATTTTTCGTTG 53184

Qy       3886 TGTGTGTATATATATATAATTATGTACTTCTGGCAATTCTATCTGTATTTAAAGATGTGA 3945
             |  || | ||| | | ||  | | |  ||||  ||       ||| | |  |  | | |
Db      53185 TTGGTTTTTATCTGTTTAGATTTCTCATTCTGAAAAAAAAGACTGAAATATAGCAGATTA 53244

Qy       3946 CAATCTTGACACCAATTTTAAGAATAGCTGTGAGACCGAATTAAAGATAATCCCTACCAA 4005
             | |  |||    |||  |   || | |       |  ||||    |||| | ||  |
Db      53245 TATTTGTGAATGAAATGCTGTTAAAATCCAGATCTCTCAATTTTTAATAACCTCTTTCGC 53304

Qy       4006 GTGAAAATTGATGTGTGTTAAGAGGGTACAGAATTATCAACTGATTTGGTCAGTTGCTTC 4065
             .|  ||    || | | |||       | |||  ||  |   || |||| |||  |  ||
Db      53305 CTCCAAGAGAATCTATTTTATAGCCTTCCAGCCTTCCCCTTTGCTTTGATCAACTAGCTC 53364


RESULT 9
US-09-949-016-54020/c
; Sequence 54020, Application US/09949016
; Patent No. 6812339
; GENERAL INFORMATION:
;  APPLICANT: VENTER, J. Craig et al.
;  TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES ASSOCIATED
;  TITLE OF INVENTION:  WITH HUMAN DISEASE, METHODS OF DETECTION AND USES THEREOF
;  FILE REFERENCE: CL001307
;  CURRENT APPLICATION NUMBER: US/09/949,016
;  CURRENT FILING DATE:  2000-04-14
;  PRIOR APPLICATION NUMBER: 60/241,755
;  PRIOR FILING DATE: 2000-10-20
;  PRIOR APPLICATION NUMBER: 60/237,768
;  PRIOR FILING DATE: 2000-10-03
;  PRIOR APPLICATION NUMBER: 60/231,498
;  PRIOR FILING DATE: 2000-09-08
;  NUMBER OF SEQ ID NOS: 207012
```

```
Qy        758 GAGGAAGGGGAATCTGATGATGAGTTTGACATGGATGAGAATCTGCCTCCCAAGTTGAGC 817
              ::::::::::::   :: :: :::   :: : ::: :::::   :      :::  :::
Db       1417 RRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRR 1358

Qy        818 AGGTTAAAGATGAATATAGCTTCTCCAGGTACAGTTCACAAACGCTACCACCGGAGGAAA 877
              :::  ::::: ::: : ::       ::: : ::   : ::: :  :  :  :::::::::
Db       1357 RRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRR 1298

Qy        878 AGTCAGGGCCGGGGCTCCAGCTGCAGTAGTTCGGAGACCAGTGATGATGATTCTGAAAGC 937
              ::  :::: ::::    ::  : :: ::   ::::: :: :: :: ::    :::::
Db       1297 RRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRR 1238

Qy        938 CGGCGGCGGCTCGATAAAGATAGCGGGTTCACCTACTCCTGGCACCGACGGGATAGCAGC 997
               :: :: ::   :: ::::: :: :::   :   :     :: :  :: :::: :: ::
Db       1237 RRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRR 1178

Qy        998 GAGGGGCCCCCTGGCAGTGAGGGGGATGGCGGGGGCCAGAGCAAGCCGAGCAATGCCAGT 1057
              ::::::      :: :: :::::::: :: :::::  :::: :::  ::: :: :  ::
Db       1177 RRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRR 1118

Qy       1058 GGAGGGGTGGACAAGGCCAGCCCCAGTGAGAACAATGCTGGTGGGGGCAGTCCCTCCAGC 1117
              ::::::: ::: ::::  ::    :: ::::: :: :  :: ::::: ::   | | |||
Db       1117 RRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRATCGCAAGC 1058


RESULT 4
US-09-949-016-14876
; Sequence 14876, Application US/09949016
; Patent No. 6812339
; GENERAL INFORMATION:
;  APPLICANT: VENTER, J. Craig et al.
;  TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES ASSOCIATED
;  TITLE OF INVENTION:  WITH HUMAN DISEASE, METHODS OF DETECTION AND USES THEREOF
;  FILE REFERENCE: CL001307
;  CURRENT APPLICATION NUMBER: US/09/949,016
;  CURRENT FILING DATE:  2000-04-14
;  PRIOR APPLICATION NUMBER: 60/241,755
;  PRIOR FILING DATE: 2000-10-20
;  PRIOR APPLICATION NUMBER: 60/237,768
;  PRIOR FILING DATE: 2000-10-03
;  PRIOR APPLICATION NUMBER: 60/231,498
;  PRIOR FILING DATE: 2000-09-08
;  NUMBER OF SEQ ID NOS: 207012
;  SOFTWARE: FastSEQ for Windows Version 4.0
; SEQ ID NO 14876
;   LENGTH: 16573
;   TYPE: DNA
;   ORGANISM: Human
;   FEATURE:
;   NAME/KEY: misc_feature
;   LOCATION: (1)...(16573)
;   OTHER INFORMATION: n = A,T,C or G
US-09-949-016-14876

  Query Match             1.2%;  Score 49.8;  DB 4;  Length 16573;
  Best Local Similarity   48.8%;  Pred. No. 0.017;
  Matches  102;  Conservative    0;  Mismatches  107;  Indels    0;  Gaps    0;

Qy       3735 AAAAACCATTACATTAAGATAATATTGGACCAAACTACTTACTTGCTCTAAACAGTTACT 3794
              |||||  || | |´ ||    ||||      ||  ||  ||  |  | |||| |  ||
Db       9766 AAAAAATATAAAAAAAAATATATATATATAATATATATATTATTTAAATATATATA 9825

Qy       3795 TGTACCCCTTAACCTGTCTTCAAAAGTTGCATATAGTTACAGTAGTGTATAAATTAAATA 3854
              | |         |  | | || ||||  |  |||||  |  |    | |||| ||| || |
Db       9826 TATTTAAAAATATATATATTAAAAATATATATATATATTTAAATATATATATATTTAAAA 9885

Qy       3855 TTGTGGAAAAACAGTCTTGTATTTTTCTGTATGTGTGTATATATATATAATTATGTACTT 3914
              | | |                 | ||| | | |||||||||  |  |||| ||  |
Db       9886 ATATATANNNNNNNNNNNNNNNNNNTATATATATTTATATATATTTTTTTATATATAT 9945

Qy       3915 CTGGCAATTCTATCTGTATTTAAAGATGT 3943
              |    | | ||| | | |||| | || |
Db       9946 ATATTTAATATATATATTTTTATATATAT 9974


RESULT 5
US-09-806-708B-22/c
; Sequence 22, Application US/09806708B
; Patent No. 6784342
; GENERAL INFORMATION:
;  APPLICANT: The University of British Columbia
;  TITLE OF INVENTION: Regulation of Embryonic Transcription in Plants
;  FILE REFERENCE: 4810-58741
;  CURRENT APPLICATION NUMBER: US/09/806,708B
;  CURRENT FILING DATE:  2001-04-03
;  PRIOR APPLICATION NUMBER: US 60/147,133
;  PRIOR FILING DATE: 1999-08-04
;  NUMBER OF SEQ ID NOS: 23
;  SOFTWARE: PatentIn version 3.0
; SEQ ID NO 22
;   LENGTH: 1141
;   TYPE: DNA
;   ORGANISM: Artificial sequence
;   FEATURE:
;   NAME/KEY: promoter
;   LOCATION: (1)..(1141)
;   OTHER INFORMATION: consensus sequence of A.t., L.a., and B.n. FAE1 promoters
US-09-806-708B-22

  Query Match             1.1%;  Score 46;  DB 4;  Length 1141;
  Best Local Similarity   10.1%;  Pred. No. 0.03;
  Matches   50;  Conservative  206;  Mismatches  238;  Indels    1;  Gaps    1;

Qy       3646 TGAGTGGTGTCTTACTCCTTTGTTTATAAGTGAATGATTGTGCATGTTTTGTATGTCATA 3705
              |                  :  | |  :|:: |   :|:| :::: :::||  ::::: : :
Db        655 TNNNNNNNNNNNNNNNNSCCTCTRMMTMRWTMKGDGMTVRKKVKWRDTTCTYVDVWADSW 596

Qy       3706 GTATGTCGTCACATAAAAGGGAGGGAGCGAAAAACCATTACATTAAGATAATATTGGACC 3765
              ::::     |: :|  :    :: | : : :   ::::    :    :| :||:
Db        595 VWWYANWMRCRDVTYTRNNTYCKSYAHSYWYWSNNAMWYRRYSARNWSSMARWTTRNNWW 536

Qy       3766 AAACTACTTACTTGCTCTAAACAGTTACTTGTACCCCTTAACCTGTCTTCAAAAGTTGCA 3825
              :  : :: :| |    ::     :|  ::: ::   :|| ::      :::::   :: :
Db        535 MSGBVRMRWAGTMWWRHWNNNNTDTRYYWWWKRWARBTTTVYDSMCNAKSMWRGNNWRAM 476

Qy       3826 TATAGTTACAGTAGTGTATAAATTAAATATTGTGGAAAAACAGTCTTGTATTTTTCTGTA 3885
              ::::     || ::: | ::   |::: : ::   :|: : | |          ::: ::
Db        475 KMWWAANNDAGAMDHWTYWMGNNTMWMRRAWKMMNMAWCRRAYCCNNNNNRACVWHKHKM 416

Qy       3886 TGTGTGTATATATATATAATTATGTACTTCTGGCAATTCTATCTGTATTTAAAGATGTGA 3945
              ::: :::: ::|      : ::  : : ::::    | ::  |:::::::  :::   :
Db        415 WRWTWKYM-WKAACNNNNBKAMYMRVAWMMYSRDTTNTDWMMWTSDWBWHWYTVDYTMMR 357

Qy       3946 CAATCTTGACACCAATTTTAAGAATAGCTGTGAGACCGAATTAAAGATAATCCCTACCAA 4005
               :        ::|  | : :: ::   :  :|     :| : ||:::||  : :: : :
Db        356 AWNNNNNNNWRBCKTTSWMWWMDHMNTHCTYGNNTWGSAYBMAAMSMWAAGASNBVTYNW 297

Qy       4006 GTGAAAATTGATGTGTGTTAAGAGGGTACAGAATTATCAACTGATTTGGTCAGTTGCTTC 4065
               :::  : : :|      :|:  : :|: :     : :  |::  :   ::|  :: ::::
Db        296 CWRMTYMGKTMTNNNNNNKAWYYRTKTVAWCNNRYYYDTAVWTBKRNYKYCYAYBWYYBM 237

Qy       4066 CAATGCTGGTTGATTTCCCTCATTGTGTAAACATTGACAGGTATGTGACAAATGGGAAAA 4125
              :: : ::: : :|:: :    : :   : ::::: |:: ::: : : |   :::::|::
Db        236 YMGKHHWBWWRRABHRSWNMWWVKCRNKYMVSWHYHAMRYBKWABAVGCNNNWKDRMAHH 177

Qy       4126 AAAATCCAAATAATA 4140
              :: ||   ::::::|
Db        176 HWCATNNNMMWWWYA 162


RESULT 6
US-09-949-016-61868/c
; Sequence 61868, Application US/09949016
; Patent No. 6812339
; GENERAL INFORMATION:
;  APPLICANT: VENTER, J. Craig et al.
```

```
Db        449 KWKANNCKWRAWDHKTCTHNNTT--WWKMKTYWNNCYWKSMTNGKSHRBAAAVYTWYMWW 506

Qy       3699 TGTCATAGTATGTCGTCACATAAAAGGGAGGGAGCGAAAAACCATTACATTAAGATAATA 3758
              :: :|:|     :: : || :   : :       :   : :|   : :     : :
Db        507 WRRYAHANNNNWDYWWKACTWYKYBVCSKWWNNYAAWYTKSSWNYTSRYYRWKTNNSWRW 566

Qy       3759 TTGGACCAAACTACTTACTTGCTCTAAACAGTTACTTGTACCCCTTAACCTGTCTTCAAA 3818
               : ::: :|       ::   : |:::::::|:::::    :::  :::: :|: : |:
Db        567 RSDTRSMGRANNYARABHYGYKWNTRWWBWSHTWBHBRAGAAHYWMBMMYBAKCHCMKAW 626

Qy       3819 AGTTGCATATAGTTACAGTAGTGTATAAATTAAATATTGTGGAAAAACAGTCTTGTATTT 3878
              : :::| |  :                  || |::: :   :  |:|    ::::: |
Db        627 YKAKKYAGAGGSNNNNNNNNNNNNNNNATCARDDYYAASRWYAMANAKWYYYKBAANNA 686

Qy       3879 TTCTGTATGTGTGTATATATATATAATTATGTACTTCTGGCAATTCTATCTGTATTTAAA 3938
              :: :   :   :  ||:|:   ::      :       |  : |: :  |  :||
Db        687 YYTHANNWWGCWNNATDTRRTMWKNNNNNNAGTWKNNNNNNAKNASAAKNYAAAAVKAAK 746

Qy       3939 GATGTGACAATCTTGACACCAATTTTAAGAATAGCTGTGAGACCGAATTAAAGATAATCC 3998
              :::::   :|   : | |     |::   || |   |       : |::|  ||:: :|
Db        747 KHWRWANKWAMRGWHADAAABTTDKRNNGAYTKYTTTNNNNTYRGVVTNTAARDGWANNN 806

Qy       3999 CTACCAAGTGAAAATTGATGTGTGTTAAGAGGGTACAGAATTATCAACTGATTTGGTCAG 4058
                      : :  :| : :  :             |  ::|  :::  ::  ::
Db        807 NNNNNNNNNNNGWSDMWVTWWAYANYGTNNNNNNNNNNAYAWWTNKWYYTTDDRWRBAYT 866

Qy       4059 TTGCTTCCAATGCTGGTTGATTTCCCTCATTGTGTAAACATTGACAGGTATGTGACAAAT 4118
                   :| :| : ::: :    |: : : : | :  |||      ::|      :
Db        867 NNNNNNRMAYYGAYADDYAYYMSDTCDAWMKWDATKMNNATTYNRGTAWRTNNNNNNMTM 926

Qy       4119 GGGAAAAAAAAATCCAAATAATAAAGTGACATATTGG 4154
              : :   :||:       :: ||:  ::  || ||:|
Db        927 KTKYYBHAAWNNNNNNGKMCTAHTWWVCKATKTTKG 962


RESULT 2
US-09-949-016-15061
; Sequence 15061, Application US/09949016
; Patent No. 6812339
; GENERAL INFORMATION:
;  APPLICANT: VENTER, J. Craig et al.
;  TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES ASSOCIATED
;  TITLE OF INVENTION:  WITH HUMAN DISEASE, METHODS OF DETECTION AND USES THEREOF
;  FILE REFERENCE: CL001307
;  CURRENT APPLICATION NUMBER: US/09/949,016
;  CURRENT FILING DATE:  2000-04-14
;  PRIOR APPLICATION NUMBER: 60/241,755
;  PRIOR FILING DATE: 2000-10-20
;  PRIOR APPLICATION NUMBER: 60/237,768
;  PRIOR FILING DATE: 2000-10-03
;  PRIOR APPLICATION NUMBER: 60/231,498
;  PRIOR FILING DATE: 2000-09-08
;  NUMBER OF SEQ ID NOS: 207012
;  SOFTWARE: FastSEQ for Windows Version 4.0
; SEQ ID NO 15061
;   LENGTH: 17612
;   TYPE: DNA
;   ORGANISM: Human
US-09-949-016-15061

  Query Match             1.3%;  Score 52.4;  DB 4;  Length 17612;
  Best Local Similarity   45.6%;  Pred. No. 0.0031;
  Matches  185;  Conservative    0;  Mismatches  221;  Indels    0;  Gaps    0;

Qy       3511 TTCAATGGTTTTTTATATTCAGATGTATATATGGTGCTCACTTTAGGATCAGCAGTGTTG 3570
              ||  ||    | | || ||  | || ||||||    | |   | || |   |      ||
Db       4916 TTTTATTTATATGTAAATGTATATTTATATAAATAGATTTATATATGTATATTTTATTTA 4975

Qy       3571 ACCATTTATGCTGCATAGCTGTATTATAGCCTTATTAGTTGTGTGGTTGACCCTTGGGGT 3630
                ||||||  |  |||  |||||  |    ||||  ||    |  |  |  |    |
Db       4976 TATATTTATATTTTATATATGTATATTTTATTTATATGTATATTTATATATAATATACAT 5035

Qy       3631 ATACAAATGTCAGTCTGAGTGGTGTCTTACTCCTTTGTTTATAAGTGAATGATTGTGCAT 3690
                ||| |||| |  || | |         |     |  | || |  |  ||   | |  ||
Db       5036 ATATAAATATACGTATTATATAATATGTATATAAATATACATATATATACATTTATAT 5095

Qy       3691 GTTTTGTATGTCATAGTATGTCGTCACATAAAAGGGAGGGAGCGAAAAACCATTACATTA 3750
                | |  ||  | | | |||       |  |||  |    |         |||| ||||| ||
Db       5096 ATATAATAAATAAAAATATACATATATTTAATACATATATTTAATAAAAACATTAAATAC 5155

Qy       3751 AGATAATATTGGACCAAACTACTTACTTGCTCTAAACAGTTACTTGTACCCCTTAACCTG 3810
              | ||| |     |  | |  | |||  |    || |||  || || ||        | |
Db       5156 ATATATTTAATAAAAATATAATTTATATATAATATACATATATTTATATATAAATATATT 5215

Qy       3811 TCTTCAAAAGTTGCATATAGTTACAGTAGTGTATAAATTAAATATTGTGGAAAAACAGTC 3870
              | |  | ||  |||||||  |||   ||        || || || ||       |||
Db       5216 TATATATAATATGCATATGTTTATTTTACATATATAAATATATTTATATATAAATATACG 5275

Qy       3871 TTGTATTTTTCTGTATGTGTGTATATATATATAATTATGTACTTCT 3916
              |    || || | ||| | | | | | | | | ||  ||| ||  | |
Db       5276 TATATTTATTTTATATATATATGTGTGTGTGTATATATATATATAT 5321


RESULT 3
US-08-232-463-14/c
; Sequence 14, Application US/08232463
; Patent No. 5670367
;  GENERAL INFORMATION:
;    APPLICANT:  DORNER, F.
;    APPLICANT:  SCHEIFLINGER, F.
;    APPLICANT:  FALKNER, F. G.
;    TITLE OF INVENTION:  RECOMBINANT FOWLPOX VIRUS
;    NUMBER OF SEQUENCES:  52
;    CORRESPONDENCE ADDRESS:
;      ADDRESSEE:  Foley & Lardner
;      STREET:  1800 Diagonal Road, Suite 500
;      CITY:  Alexandria
;      STATE:  VA
;      COUNTRY:  USA
;      ZIP:  22313-0299
;    COMPUTER READABLE FORM:
;      MEDIUM TYPE:  Floppy disk
;      COMPUTER:  IBM PC compatible
;      OPERATING SYSTEM:  PC-DOS/MS-DOS
;      SOFTWARE:  PatentIn Release #1.0, Version #1.25
;    CURRENT APPLICATION DATA:
;      APPLICATION NUMBER:  US/08/232,463
;      FILING DATE:
;      CLASSIFICATION:  435
;    PRIOR APPLICATION DATA:
;      APPLICATION NUMBER:  US/07/935,313
;      FILING DATE:
;      APPLICATION NUMBER:  EP 91 114 300.6
;      FILING DATE:  26-AUG-1991
;    ATTORNEY/AGENT INFORMATION:
;      NAME:  BENT, Stephen A.
;      REGISTRATION NUMBER:  29,768
;      REFERENCE/DOCKET NUMBER:  30472/114 IMMU
;    TELECOMMUNICATION INFORMATION:
;      TELEPHONE:  (703)836-9300
;      TELEFAX:  (703)683-4109
;      TELEX:  899149
;  INFORMATION FOR SEQ ID NO:  14:
;    SEQUENCE CHARACTERISTICS:
;      LENGTH:  7218 base pairs
;      TYPE:  nucleic acid
;      STRANDEDNESS:  single
;      TOPOLOGY:  linear
;    IMMEDIATE SOURCE:
;      CLONE:  pTZgpt-F1s
US-08-232-463-14

  Query Match             1.2%;  Score 50;  DB 1;  Length 7218;
  Best Local Similarity   1.4%;  Pred. No. 0.008;
  Matches    5;  Conservative  215;  Mismatches  140;  Indels    0;  Gaps    0;
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Post-processing: Minimum Match 0%
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Listing first 45 summaries

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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

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1	56.2	1.3	1141	4	US-09-806-708B-22	Sequence 22, Appl
2	52.4	1.3	17612	4	US-09-949-016-15061	Sequence 15061, A
c 3	50	1.2	7218	1	US-08-232-463-14	Sequence 14, Appl
4	49.8	1.2	16573	4	US-09-949-016-14876	Sequence 14876, A
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c 7	45.4	1.1	314798	4	US-09-949-016-13539	Sequence 13539, A
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c 45	42	1.0	134987	4	US-09-949-016-15349	Sequence 15349, A

ALIGNMENTS

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; Sequence 22, Application US/09806708B
; Patent No. 6784342
; GENERAL INFORMATION:
; APPLICANT: The University of British Columbia
; TITLE OF INVENTION: Regulation of Embryonic Transcription in Plants
; FILE REFERENCE: 4810-58741
; CURRENT APPLICATION NUMBER: US/09/806,708B
; CURRENT FILING DATE: 2001-04-03
; PRIOR APPLICATION NUMBER: US 60/147,133
; PRIOR FILING DATE: 1999-08-04
; NUMBER OF SEQ ID NOS: 23
; SOFTWARE: PatentIn version 3.0
; SEQ ID NO 22
; LENGTH: 1141
; TYPE: DNA
; ORGANISM: Artificial sequence
; FEATURE:
; NAME/KEY: promoter
; LOCATION: (1)..(1141)
; OTHER INFORMATION: consensus sequence of A.t., L.a., and B.n. FAE1 promoters
US-09-806-708B-22

Query Match 1.3%; Score 56.2; DB 4; Length 1141;
Best Local Similarity 11.8%; Pred. No. 3.3e-05;
Matches 96; Conservative 285; Mismatches 433; Indels 2; Gaps 1;

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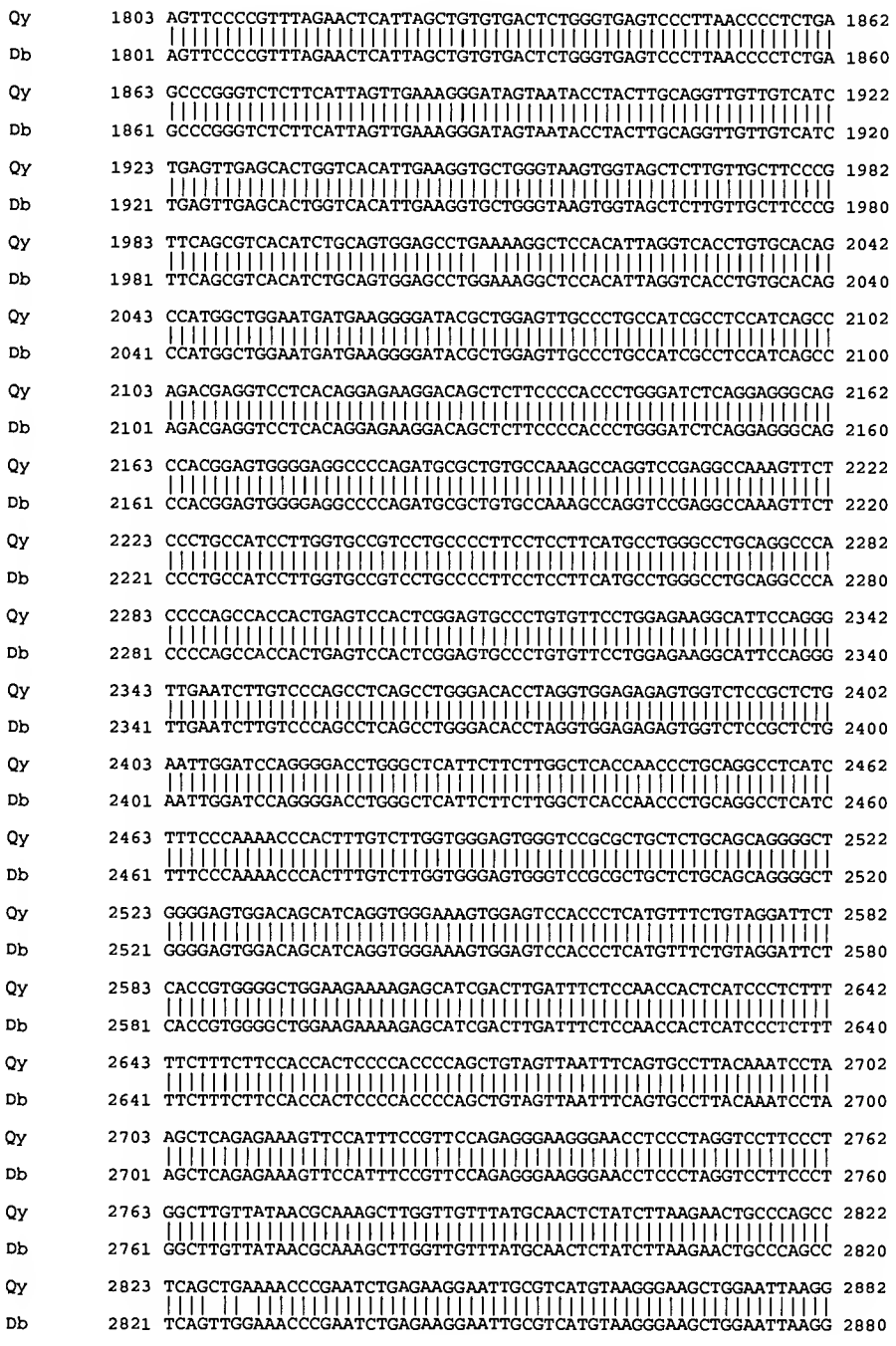
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;  PRIOR APPLICATION NUMBER: US/09/997,263
;  PRIOR FILING DATE: 2001-11-30
;  PRIOR APPLICATION NUMBER: US 09/802,806
;  PRIOR FILING DATE: 2001-03-09
;  PRIOR APPLICATION NUMBER: US 60/207,333
;  PRIOR FILING DATE: 2000-05-30
;  PRIOR APPLICATION NUMBER: US 09/384,446
;  PRIOR FILING DATE: 1999-08-27
;  PRIOR APPLICATION NUMBER: US 60/132,684
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US-10-857-942-3

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Qy        903 GAGCCCAGCAAGCTGAATAGCATTGGCAATTACTACAAGCCGTGGTTCTTTAAGCATGTG 962
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Qy        963 GAGAACTATCTGAAGACAAACCGAGAGGGCCTGGAGTACATTCCCTTGAGACACTACTAC 1022
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Db        961 GAGAACTATCTGAAGACAAACCGAGAGGGCCTGGAGTACATTCCCTTGAGACACTACTAC 1020

Qy       1023 CACCGCCACACGCGCAGCATCTTCTGGGAGCTCCAGGACATCATCCCCTTTGGCAACAAC 1082
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Qy       1083 CCCATCTTCCGCTACCTCTTTGGCTGGATGGTGCCTCCCAAGATCTCCCTCCTGAAGCTG 1142
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Db       1081 CCCATCTTCCGCTACCTCTTTGGCTGGATGGTGCCTCCCAAGATCTCCCTCCTGAAGCTG 1140

Qy       1143 ACCCAGGGTGAGACCCTGCGCAAGCTGTACGAGCAGCACCACGTGGTGCAGGACATGCTG 1202
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Qy       1203 GTGCCCATGAAGTGCCTGCAGCAGGCCCTGCACACCTTCCAAAACGACATCCACGTCTAC 1262
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Qy       1323 AATGAGGCAGAGCTCTACATCGACATTGGAGCATATGGGGAGCCGCGTGTGAAACACTTT 1382
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Db       1321 AATGAGGCAGAGCTCTACATCGACATTGGAGCATATGGGGAGCCGCGTGTGAAACACTTT 1380

Qy       1383 GAAGCCAGGTCCTGCATGAGGCAGCTGGAGAAGTTTGTCCGCAGCGTGCATGGCTTCCAG 1442
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Db       1381 GAAGCCAGGTCCTGCATGAGGCAGCTGGAGAAGTTTGTCCGCAGCGTGCATGGCTTCCAG 1440

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Qy       1743 AGGGGGCAGCCCAGTGGAATGGAAAGAATGTGGGATTTGGAGTCAGACAAGCCTGAGTCC 1802
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Db       1741 AGGGGGCAGCCCAGTGGAATGGAAAGAATGTGGGATTTGGAGTCAGACAAGCCTGAGTCC 1800
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Db       2264 CACCCCAGCCACCACTGAGTCCACTCGGAGTGCCCTGTGTTCCTGGAGAAGGCATTCCAG 2323

Qy       2341 GGTTGAATCTTGTCCCAGCCTCAGCCTGGGACACCTAGGTGGAGAGAGTGGTCTCCGCTC 2400
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Db       2324 GGTTGAATCTTGTCCCAGCCTCAGCCTGGGACACCTAGGTGGAGAGAGTGGTCTCCGCTC 2383

Qy       2401 TGAATTGGATCCAGGGGACCTGGGCTCATTCTTCTTGGCTCACCAACCCTGCAGGCCTCA 2460
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Qy       2461 TCTTTCCCAAAACCCACTTTGTCTTGGTGGGAGTGGGTCCGCGCTGCTCTGCAGCAGGGG 2520
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Qy       2641 TTTTCTTTCTTCCACCACTCCCCACCCCAGCTGTAGTTAATTTCAGTGCCTTACAAATCC 2700
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Qy       2761 CTGGCTTGTTATAACGCAAAGCTTGGTTGTTTATGCAACTCTATCTTAAGAACTGCCCAG 2820
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Db       3644 CCACAACCCGGGGGCTTGGAAAGGAAGGTATTTTGGAATCACACCCTCCGGTTATGTTGCT 3703

Qy       3721 CCAGTAAAATCTTGCCTGGAAAGAGGCAGTCTTCTTAGCATGGTGAGCTGAGTTCATGGC 3780
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Db       3764 TTTTTTTTGTAGCCAGTCCTGTCCCTGGCCATCCATGTGATGGTTTTGGATGGAGTTAAA 3823

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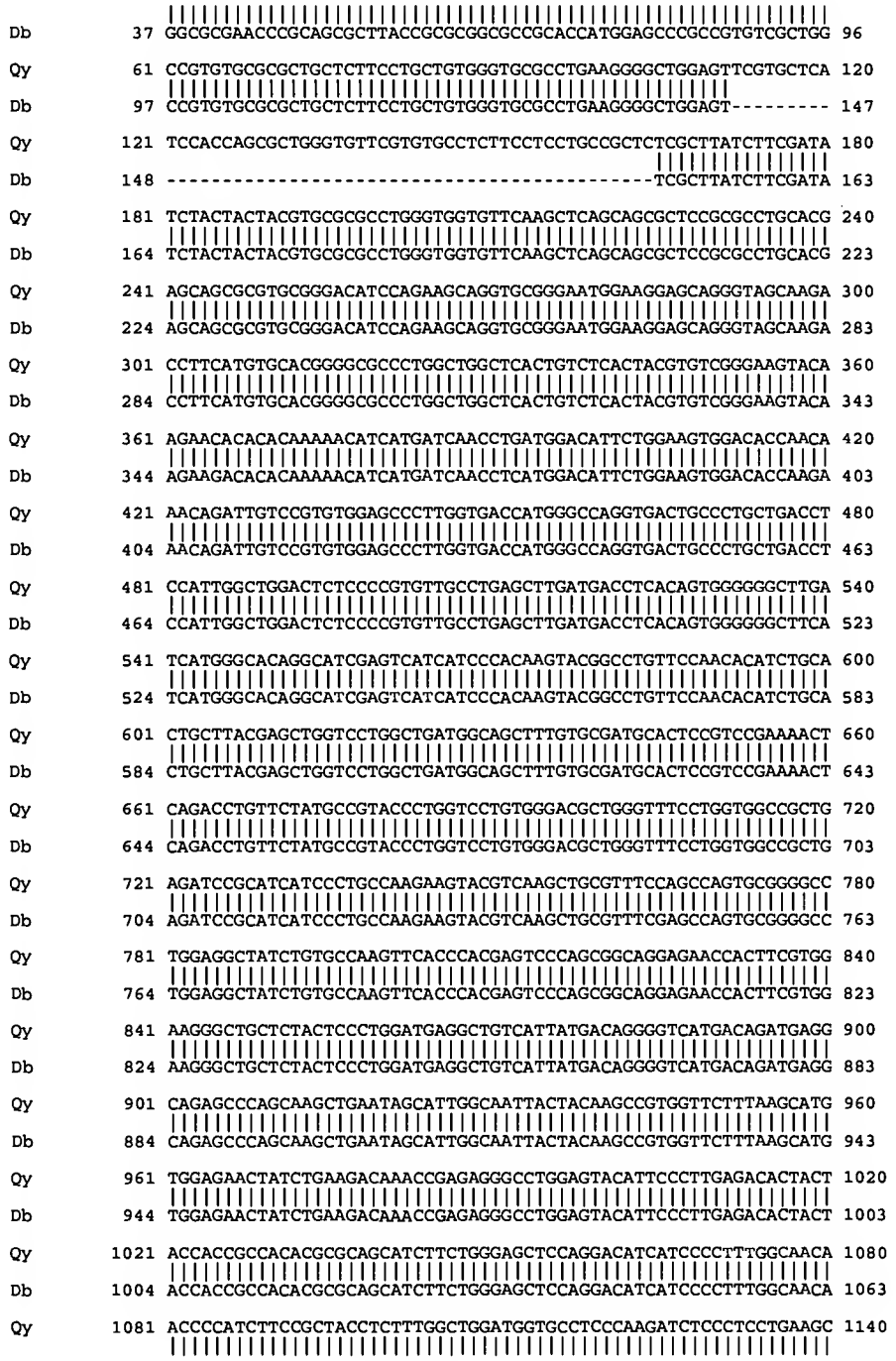
RESULT 15
US-10-857-942-3
; Sequence 3, Application US/10857942
; Publication No. US20050004065A1
; GENERAL INFORMATION:
; APPLICANT: EINAT, Paz
; APPLICANT: FEINSTEIN, Elena
; APPLICANT: SKALITER, Rami
; TITLE OF INVENTION: HYPOXIA REGULATED GENES
; FILE REFERENCE: FEINSTEIN=5.1B
; CURRENT APPLICATION NUMBER: US/10/857,942
; CURRENT FILING DATE: 2004-06-02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Qy       2641 TTTTCTTTCTTCCACCACTCCCCACCCCAGCTGTAGTTAATTTCAGTGCCTTACAAATCC 2700
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Db       2702 TTTTCTTTCTTCCACCACTCCCCACCCCAGCTGTAGTTAATTTCAGTGCCTTACAAATCC 2761

Qy       2701 TAAGCTCAGAGAAAGTTCCATTTCCGTTCCAGAGGGAAGGGAACCTCCCTAGGTCCTTCC 2760
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Db       2762 TAAGCTCAGAGAAAGTTCCATTTCCGTTCCAGAGGGAAGGGAACCTCCCTAGGTCCTTCC 2821

Qy       2761 CTGGCTTGTTATAACGCAAAGCTTGGTTGTTTATGCAACTCTATCTTAAGAACTGCCCAG 2820
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Db       2822 CTGGCTTGTTATAACGCAAAGCTTGGTTGTTTATGCAACTCTATCTTAAGAACTGCCCAG 2881

Qy       2821 CCTCAGCTGAAAACCCGAATCTGAGAAGGAATTGCGTCATGTAAGGGAAGCTGGAATTAA 2880
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       2882 CCTCAGCTGAAAACCCGAATCTGAGAAGGAATTGCGTCATGTAAGGGAAGCTGGAATTAA 2941

Qy       2881 GGGAGCTGAGCCAGTCATGGTTGTGGCGTGTGAGTCAGGAGACCTAGGTTTCAGCCCCTC 2940
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       2942 GGGAGCTGAGCCAGTCATGGTTGTGGCGTGTGAGTCAGGAGACCTAGGTTTCAGCCCCTC 3001

Qy       2941 TCTACTGTCAGCGAGCTGTGCAACGTGGGCAAGTCATTGTCCTCTGAGCTGCAGTTTCCT 3000
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       3002 TCTACTGTCAGCGAGCTGTGCAACGTGGGCAAGTCATTGTCCTCTGAGCTGCAGTTTCCT 3061

Qy       3001 CATCTGTCACATCGCTACAGACAAGACCTCCCTGGAACCCTTCTGATTGTCTTAGACACT 3060
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Qy       3061 GTGGTTGCAAAACCCACGGAAAGCCTCATTTGTGTGGAAAGTCAGAGGAAAAATGATCCA 3120
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Db       3122 GTGGTTGCAAAACCCACGGAAAGCCTCATTTGTGTGGAAAGTCAGAGGAAAAATGATCCA 3181

Qy       3121 GTGGACACTTGGGGATTATCTGTCATTCAAGATCCTTCCTTCAACCCCAAGGCCAGCTCC 3180
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Db       3182 GTGGACACTTGGGGATTATCTGTCATTCAAGATCCTTCCTTCAACCCCAAGGTCAGCTCC 3241

Qy       3181 CATCTCATTTCCAGAAAGGCTCATACCTGGCTTGCAGGGAAGCATCTGTCTTGTCATTCC 3240
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Qy       3241 AGGTGCCAGAATCCTCTCAGAGTCATTGAAGGGTGTTCACCCATCCCACCCAAGGCTTGG 3300
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Db       3302 AGGTGCCAGAATCCTCTCAGAGTCATTGAAGGGTGTTCACCCATCCCACCCAAGGCTTGG 3361

Qy       3301 CACACTGCCAGTGTCTTAGCAGGGTCTTGTGAGGGCTGGGGGCATCCAGGCACTCAGAAG 3360
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Qy       3361 GCAAAGGAACCACCCTACCCATTTGGCCTCTGGAGGGGGCAGAAGAAAGAAAGAAACCTC 3420
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Db       3422 GCAAAGGAACCACCCTACCCATTTGGCCTCTGGAGGGGGCAGAAGAAAGAAAGAAACCTC 3481

Qy       3421 ATCCTATATTTTACAAAGCATGTGAATTCTGGCATTAGCTCTCATAGGAGACCCATGTGC 3480
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Db       3482 ATCCTATATTTTACAAAGCATGTGAATTCTGGCATTAGCTCTCATAGGAGACCCATGTGC 3541

Qy       3481 TTCCTTGCTCAGTGCAAAACTGATGATTCTACTTGCTGTAGATGAATGGTTAACACGAGC 3540
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Db       3542 TTCCTTGCTCAGTGCAAAACTGATGATTCTACTTGCTGTAGATGAATGGTTAACACGAGC 3601

Qy       3541 TAGTTAAACAGTGCCATTGTTTTGCCAGTGAAGCCTCCAACCCTAAGCCACTGGGACGGT 3600
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Db       3602 TAGTTAAACAGTGCCATTGTTTTGCCAGTGAAGCCTCCAACCCTAAGCCACTGGGACGGT 3661

Qy       3601 GGCCAGAGATGCCAGCAGCCTCTGTCGCCCTTAGTCATATAACCAAAATCCAGACCTTAT 3660
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       3662 GGCCAGAGATGCCAGCAGCCTCTGTCGCCCTTAGTCATATAACCAAAATCCAGACCTTAT 3721

Qy       3661 CCACAACCCGGGGCTTGGAAAGGAAGGTATTTTGGAATCACACCCTCCGGTTATGTTGCT 3720
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       3722 CCACAACCCGGGGCTTGGAAAGGAAGGTATTTTGGAATCACACCCTCCGGTTATGTTGCT 3781

Qy       3721 CCAGTAAAATCTTGCCTGGAAAGAGGCAGTCTTCTTAGCATGGTGAGCTGAGTTCATGGC 3780
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       3782 CCAGTAAAATCTTGCCTGGAAAGAGGCAGTCTTCTTAGCATGGTGAGCTGAGTTCATGGC 3841

Qy       3781 TTTTTTTTGTAGCCAGTCCTGTCCCTGGCCATCCATGTGATGGTTTTGGATGGAGTTAAA 3840
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       3842 TTTTTTTTGTAGCCAGTCCTGTCCCTGGCCATCCATGTGATGGTTTTGGATGGAGTTAAA 3901

Qy       3841 CTTGATGCCAGTGGGCAGTGCATGTGGAAAGTATCAGAGTAAGCCTCTCCCCTCCAGAGC 3900
             |||||||||||||||||||||||||||||||||||||||||||||| ||||||||||||||||
Db       3902 CTTGATGCCAGTGGGCAGTGCATGTGGAAAGTATCAGAGTAAGGCTCTCCCCTCCAGAGC 3961

Qy       3901 CCTGAGTTTCTTGGCTGCATGAAGGTTTTCTTTAGAATCAGAATTGTAGCCAGTTTCTTT 3960
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Db       3962 CCTGAGTTTCTTGGCTGCATGAAGGTTTTCTTTAGAATCAGAATTGTAGCCAGTTTCTTT 4021

Qy       3961 GGCCAGAAGGATGAATACTTGGATATTACTGAAAGGGAGGGGTGGAGATGGGTGTGGCAG 4020
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Db       4022 GGCCAGAAGGATGAATACTTGGATATTACTGAAAGGGAGGGGTGGAGATGGGTGTGGCAG 4081

Qy       4021 TGTATGGTGTGTGATTTTTATTTTCTTCTTTGGTCATGGGGGCCAAGGAGAAAGGCATGA 4080
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Qy       4081 ATCTTCCCTGTCAGGCTCTTACAGCCACAGGCACTGTGTCTACTGTCTGGAAGACATGTC 4140
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Db       4142 ATCTTCCCTGTCAGGCTCTTACAGCCACAGGCACTGTGTCTACTGTCTGGAAGACATGTC 4201

Qy       4141 CCCGTGGCTGTGGGGCCGCTGCTTCTGTTTAAATAAAAGTGGCCTGG 4187
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Db       4202 CCCGTGGCTGTGGGGCCGCTGCTTCTGTTTAAATAAAAGTGGCCTGG 4248
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RESULT 14
US-10-479-081-121
; Sequence 121, Application US/10479081
; Publication No. US20050059001A1
; GENERAL INFORMATION:
; APPLICANT: NAKAGAWARA, AKIRA
; TITLE OF INVENTION: NUCLEIC ACIDS ISOLATED FROM NEUROBLASTOMA
; FILE REFERENCE: 7388-80893
; CURRENT APPLICATION NUMBER: US/10/479,081
; CURRENT FILING DATE: 2003-11-26
; PRIOR APPLICATION NUMBER: PCT/JP02/05295
; PRIOR FILING DATE: 2002-05-30
; PRIOR APPLICATION NUMBER: JP 2001-163666
; PRIOR FILING DATE: 2001-05-31
; PRIOR APPLICATION NUMBER: JP 2001-255260
; PRIOR FILING DATE: 2001-08-24
; NUMBER OF SEQ ID NOS: 742
; SOFTWARE: PatentIn version 3.2
; SEQ ID NO 121
; LENGTH: 4219
; TYPE: DNA
; ORGANISM: Homo sapiens
; FEATURE:
; OTHER INFORMATION: nbla-03646
US-10-479-081-121

Query Match 97.2%; Score 4067.8; DB 21; Length 4219;
Best Local Similarity 98.7%; Pred. No. 0;
Matches 4132; Conservative 0; Mismatches 2; Indels 53; Gaps 1;

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Qy          1 GGCGCGAACCCGCAGCGCTTACCGCGCGGCGCCGCACCATGGAGCCCGCCGTGTCGCTGG 60
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Qy       301 CCTTCATGTGCACGGGGCGCCCTGGCTGGCTCACTGTCTCACTACGTGTCGGGAAGTACA 360
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Qy       361 AGAAGACACACAAAAACATCATGATCAACCTGATGGACATTCTGGAAGTGGACACCAAGA 420
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Qy       421 AACAGATTGTCCGTGTGGAGCCCTTGGTGACCATGGGCCAGGTGACTGCCCTGCTGACCT 480
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Qy       481 CCATTGGCTGGACTCTCCCCGTGTTGCCTGAGCTTGATGACCTCACAGTGGGGGGCTTGA 540
Db       542 CCATTGGCTGGACTCTCCCCGTGTTGCCTGAGCTTGATGACCTCACAGTGGGGGGCTTGA 601

Qy       541 TCATGGGCACAGGCATCGAGTCATCATCCCACAAGTACGGCCTGTTCCAACACATCTGCA 600
Db       602 TCATGGGCACAGGCATCGAGTCATCATCCCACAAGTACGGCCTGTTCCAACACATCTGCA 661

Qy       601 CTGCTTACGAGCTGGTCCTGGCTGATGGCAGCTTTGTGCGATGCACTCCGTCCGAAAACT 660
Db       662 CTGCTTACGAGCTGGTCCTGGCTGATGGCAGCTTTGTGCGATGCACTCCGTCCGAAAACT 721

Qy       661 CAGACCTGTTCTATGCCGTACCCTGGTCCTGTGGGACGCTGGGTTTCCTGGTGGCCGCTG 720
Db       722 CAGACCTGTTCTATGCCGTACCCTGGTCCTGTGGGACGCTGGGTTTCCTGGTGGCCGCTG 781

Qy       721 AGATCCGCATCATCCCTGCCAAGAAGTACGTCAAGCTGCGTTTCGAGCCAGTGCGGGGCC 780
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Qy       781 TGGAGGCTATCTGTGCCAAGTTCACCCACGAGTCCCAGCGGCAGGAGAACCACTTCGTGG 840
Db       842 TGGAGGCTATCTGTGCCAAGTTCACCCACGAGTCCCAGCGGCAGGAGAACCACTTCGTGG 901

Qy       841 AAGGGCTGCTCTACTCCCTGGATGAGGCTGTCATTATGACAGGGGTCATGACAGATGAGG 900
Db       902 AAGGGCTGCTCTACTCCCTGGATGAGGCTGTCATTATGACAGGGGTCATGACAGATGAGG 961

Qy       901 CAGAGCCCAGCAAGCTGAATAGCATTGGCAATTACTACAAGCCGTGGTTCTTTAAGCATG 960
Db       962 CAGAGCCCAGCAAGCTGAATAGCATTGGCAATTACTACAAGCCGTGGTTCTTTAAGCATG 1021

Qy       961 TGGAGAACTATCTGAAGACAAACCGAGAGGGCCTGGAGTACATTCCCTTGAGACACTACT 1020
Db      1022 TGGAGAACTATCTGAAGACAAACCGAGAGGGCCTGGAGTACATTCCCTTGAGACACTACT 1081

Qy      1021 ACCACCGCCACACGCGCAGCATCTTCTGGGAGCTCCAGGACATCATCCCCTTTGGCAACA 1080
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Qy      1081 ACCCCATCTTCCGCTACCTCTTTGGCTGGATGGTGCCTCCCAAGATCTCCCTCCTGAAGC 1140
Db      1142 ACCCCATCTTCCGCTACCTCTTTGGCTGGATGGTGCCTCCCAAGATCTCCCTCCTGAAGC 1201

Qy      1141 TGACCCAGGGTGAGACCCTGCGCAAGCTGTACGAGCAGCACCACGTGGTGCAGGACATGC 1200
Db      1202 TGACCCAGGGTGAGACCCTGCGCAAGCTGTACGAGCAGCACCACGTGGTGCAGGACATGC 1261

Qy      1201 TGGTGCCCATGAAGTGCCTGCAGCAGGCCCTGCACACCTTCCAAAACGACATCCACGTCT 1260
Db      1262 TGGTGCCCATGAAGTGCCTGCAGCAGGCCCTGCACACCTTCCAAAACGACATCCACGTCT 1321

Qy      1261 ACCCCATCTGGCTGTGTCCGTTCATCCTGCCCAGCCAGCCAGGCCTAGTGCACCCCAAAG 1320
Db      1322 ACCCCATCTGGCTGTGTCCGTTCATCCTGCCCAGCCAGCCAGGCCTAGTGCACCCCAAAG 1381

Qy      1321 GAAATGAGGCAGAGCTCTACATCGACATTGGAGCATATGGGGAGCCGCGTGTGAAACACT 1380
Db      1382 GAAATGAGGCAGAGCTCTACATCGACATTGGAGCATATGGGGAGCCGCGTGTGAAACACT 1441

Qy      1381 TTGAAGCCAGGTCCTGCATGAGGCAGCTGGAGAAGTTTGTCCGCAGCGTGCATGGCTTCC 1440

Db      1442 TTGAAGCCAGGTCCTGCATGAGGCAGCTGGAGAAGTTTGTCCGCAGCGTGCATGGCTTCC 1501

Qy      1441 AGATGCTGTATGCCGACTGCTACATGAACCGGGAGGAGTTCTGGGAGATGTTTGATGGCT 1500
Db      1502 AGATGCTGTATGCCGACTGCTACATGAACCGGGAGGAGTTCTGGGAGATGTTTGATGGCT 1561

Qy      1501 CCTTGTACCACAAGCTGCGAGAGAAGCTGGGTTGCCAGGACGCCTTCCCCGAGGTGTACG 1560
Db      1562 CCTTGTACCACAAGCTGCGAGAGAAGCTGGGTTGCCAGGACGCCTTCCCCGAGGTGTACG 1621

Qy      1561 ACAAGATCTGCAAGGCCGCCAGGCACTGAGCTGGAGCCCGCCTGGAGAGACAGACACGTG 1620
Db      1622 ACAAGATCTGCAAGGCCGCCAGGCACTGAGCTGGAGCCCGCCTGGAGAGACAGACACGTG 1681

Qy      1621 TGAGTGGTCAGGCATCTTCCCTTCACTCAAGCTTGGCTGCTTTCCTAGATCCACACTTTC 1680
Db      1682 TGAGTGGTCAGGCATCTTCCCTTCACTCAAGCTTGGCTGCTTTCCTAGATCCACACTTTC 1741

Qy      1681 AAAGAGAAACCCCTCCAGAACTCCCACCCTGACAGCCCAACACCACCTTCCTCCTGGCTT 1740
Db      1742 AAAGAGAAACCCCTCCAGAACTCCCACCCTGACAGCCCAACACCACCTTCCTCCTGGCTT 1801

Qy      1741 CCAGGGGGCAGCCCAGTGGAATGGAAAGAATGTGGGATTTGGAGTCAGACAAGCCTGAGT 1800
Db      1802 CCAGGGGGCAGCCCAGTGGAATGGAAAGAATGTGGGATTTGGAGTCAGACAAGCCTGAGT 1861

Qy      1801 CCAGTTCCCCGTTTAGAACTCATTAGCTGTGTGACTCTGGGTGAGTCCCTTAACCCCTCT 1860
Db      1862 CCAGTTCCCCGTTTAGAACTCATTAGCTGTGTGACTCTGGGTGAGTCCCTTAACCCCTCT 1921

Qy      1861 GAGCCCGGGTCTCTTCATTAGTTGAAAGGGATAGTAATACCTACTTGCAGGTTGTTGTCA 1920
Db      1922 GAGCCCGGGTCTCTTCATTAGTTGAAAGGGATAGTAATACCTACTTGCAGGTTGTTGTCA 1981

Qy      1921 TCTGAGTTGAGCACTGGTCACATTGAAGGTGCTGGGTAAGTGGTAGCTCTTGTTGCTTCC 1980
Db      1982 TCTGAGTTGAGCACTGGTCACATTGAAGGTGCTGGGTAAGTGGTAGCTCTTGTTGCTTCC 2041

Qy      1981 CGTTCAGCGTCACATCTGCAGTGGAGCCTGAAAAGGCTCCACATTAGGTCACCTGTGCAC 2040
Db      2042 CGTTCAGCGTCACATCTGCAGTGGAGCCTGAAAAGGCTCCACATTAGGTCACCTGTGCAC 2101

Qy      2041 AGCCATGGCTGGAATGATGAAGGGGATACGCTGGAGTTGCCCTGCCATCGCCTCCATCAG 2100
Db      2102 AGCCATGGCTGGAATGATGAAGGGGATACGCTGGAGTTGCCCTGCCATCGCCTCCATCAG 2161

Qy      2101 CCAGACGAGGTCCTCACAGGAGAAGGACAGCTCTTCCCCACCCTGGGATCTCAGGAGGGC 2160
Db      2162 CCAGACGAGGTCCTCACAGGAGAAGGACAGCTCTTCCCCACCCTGGGATCTCAGGAGGGC 2221

Qy      2161 AGCCACGGAGTGGGGAGGCCCCAGATGCGCTGTGCCAAAGCCAGGTCCGAGGCCAAAGTT 2220
Db      2222 AGCCACGGAGTGGGGAGGCCCCAGATGCGCTGTGCCAAAGCCAGGTCCGAGGCCAAAGTT 2281

Qy      2221 CTCCCTGCCATCCTTGGTGCCGTCCTGCCCCTTCCTCCTTCATGCCTGGGCCTGCAGGCC 2280
Db      2282 CTCCCTGCCATCCTTGGTGCCGTCCTGCCCCTTCCTCCTTCATGCCTGGGCCTGCAGGCC 2341

Qy      2281 CACCCCAGCCACCACTGAGTCCACTCGGAGTGCCCTGTGTTCCTGGAGAAGGCATTCCAG 2340
Db      2342 CACCCCAGCCACCACTGAGTCCACTCGGAGTGCCCTGTGTTCCTGGAGAAGGCATTCCAG 2401

Qy      2341 GGTTGAATCTTGTCCCAGCCTCAGCCTGGGACACCTAGGTGGAGAGAGTGGTCTCCGCTC 2400
Db      2402 GGTTGAATCTTGTCCCAGCCTCAGCCTGGGACACCTAGGTGGAGAGAGTGGTCTCCGCTC 2461

Qy      2401 TGAATTGGATCCAGGGGACCTGGGCTCATTCTTCTTGGCTCACCAACCCTGCAGGCCTCA 2460
Db      2462 TGAATTGGATCCAGGGGACCTGGGCTCATTCTTCTTGGCTCACCAACCCTGCAGGCCTCA 2521

Qy      2461 TCTTTCCCAAAACCCACTTTGTCTTGGTGGGAGTGGGTCCGCGCTGCTCTGCAGCAGGGG 2520
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Qy       2761 CTGGCTTGTTATAACGCAAAGCTTGGTTGTTTATGCAACTCTATCTTAAGAACTGCCCAG 2820
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Db       2796 CTGGCTTGTTATAACGCAAAGCTTGGTTGTTTATGCAACTCTATCTTAAGAACTGCCCAG 2855

Qy       2821 CCTCAGCTGAAAACCCGAATCTGAGAAGGAATTGCGTCATGTAAGGGAAGCTGGAATTAA 2880
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       2856 CCTCAGCTGAAAACCCGAATCTGAGAAGGAATTGCGTCATGTAAGGGAAGCTGGAATTAA 2915

Qy       2881 GGGAGCTGAGCCAGTCATGGTTGTGGCGTGTGAGTCAGGAGACCTAGGTTTCAGCCCCTC 2940
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       2916 GGGAGCTGAGCCAGTCATGGTTGTGGCGTGTGAGTCAGGAGACCTAGGTTTCAGCCCCTC 2975

Qy       2941 TCTACTGTCAGCGAGCTGTGCAACGTGGGCAAGTCATTGTCCTCTGAGCTGCAGTTTCCT 3000
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       2976 TCTACTGTCAGCGAGCTGTGCAACGTGGGCAAGTCATTGTCCTCTGAGCTGCAGTTTCCT 3035

Qy       3001 CATCTGTCACATCGCTACAGACAAGACCTCCCTGGAACCCTTCTGATTGTCTTAGACACT 3060
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Db       3036 CATCTGTCACATCGCTACAGACAAGACCTCCCTGGAACCCTTCTGATTGTCTTAGACACT 3095

Qy       3061 GTGGTTGCAAAACCCACGGAAAGCCTCATTTGTGTGGAAAGTCAGAGGAAAAATGATCCA 3120
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Qy       3121 GTGGACACTTGGGGATTATCTGTCATTCAAGATCCTTCCTTCAACCCCAAGGCCAGCTCC 3180
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Qy       3181 CATCTCATTTCCAGAAAGGCTCATACCTGGCTTGCAGGGAAGCATCTGTCTTGTCATTCC 3240
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Qy       3241 AGGTGCCAGAATCCTCTCAGAGTCATTGAAGGGTGTTCACCCATCCCACCCAAGGCTTGG 3300
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Db       3276 AGGTGCCAGAATCCTCTCAGAGTCATTGAAGGGTGTTCACCCATCCCACCCAAGGCTTGG 3335

Qy       3301 CACACTGCCAGTGTCTTAGCAGGGTCTTGTGAGGGCTGGGGGCATCCAGGCACTCAGAAG 3360
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Qy       3361 GCAAAGGAACCACCCTACCCATTTGGCCTCTGGAGGGGGCAGAAGAAAGAAAGAAACCTC 3420
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Db       3396 GCAAAGGAACCACCCTACCCATTTGGCCTCTGGAGGGGGCAGAAGAAAGAAAGAAACCTC 3455

Qy       3421 ATCCTATATTTTACAAAGCATGTGAATTCTGGCATTAGCTCTCATAGGAGACCCATGTGC 3480
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Db       3456 ATCCTATATTTTACAAAGCATGTGAATTCTGGCATTAGCTCTCATAGGAGACCCATGTGC 3515

Qy       3481 TTCCTTGCTCAGTGCAAAACTGATGATTCTACTTGCTGTAGATGAATGGTTAACACGAGC 3540
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Db       3516 TTCCTTGCTCAGTGCAAAACTGATGATTCTACTTGCTGTAGATGAATGGTTAACACGAGC 3575

Qy       3541 TAGTTAAACAGTGCCATTGTTTTGCCAGTGAAGCCTCCAACCCTAAGCCACTGGGACGGT 3600
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Db       3576 TAGTTAAACAGTGCCATTGTTTTGCCAGTGAAGCCTCCAACCCTAAGCCACTGGGACGGT 3635

Qy       3601 GGCCAGAGATGCCAGCAGCCTCTGTCGCCCTTAGTCATATAACCAAAATCCAGACCTTAT 3660
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Qy       3661 CCACAACCCGGGGCTTGGAAAGGAAGGTATTTTGGAATCACACCCTCCGGTTATGTTGCT 3720
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Db       3696 CCACAACCCGGGGCTTGGAAAGGAAGGTATTTTGGAATCACACCCTCCGGTTATGTTGCT 3755

Qy       3721 CCAGTAAAATCTTGCCTGGAAAGAGGCAGTCTTCTTAGCATGGTGAGCTGAGTTCATGGC 3780
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Db       3756 CCAGTAAAATCTTGCCTGGAAAGAGGCAGTCTTCTTAGCATGGTGAGCTGAGTTCATGGC 3815

Qy       3781 TTTTTTTTGTAGCCAGTCCTGTCCCTGGCCATCCATGTGATGGTTTTGGATGGAGTTAAA 3840
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Db       3816 TTTTTTTTGTAGCCAGTCCTGTCCCTGGCCATCCATGTGATGGTTTTGGATGGAGTTAAA 3875
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Qy       3841 CTTGATGCCAGTGGGCAGTGCATGTGGAAAGTATCAGAGTAAGCCTCTCCCCTCCAGAGC 3900
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Db       3876 CTTGATGCCAGTGGGCAGTGCATGTGGAAAGTATCAGAGTAAGCCTCTCCCCTCCAGAGC 3935

Qy       3901 CCTGAGTTTCTTGGCTGCATGAAGGTTTTCTTTAGAATCAGAATTGTAGCCAGTTTCTTT 3960
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Db       3936 CCTGAGTTTCTTGGCTGCATGAAGGTTTTCTTTAGAATCAGAATTGTAGCCAGTTTCTTT 3995

Qy       3961 GGCCAGAAGGATGAATACTTGGATATTACTGAAAGGGAGGGGTGGAGATGGGTGTGGCAG 4020
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       3996 GGCCAGAAGGATGAATACTTGGATATTACTGAAAGGGAGGGGTGGAGATGGGTGTGGCAG 4055

Qy       4021 TGTATGGTGTGTGATTTTTATTTTCTTCTTTGGTCATGGGGGCCAAGGAGAAAGGCATGA 4080
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Qy       4081 ATCTTCCCTGTCAGGCTCTTACAGCCACAGGCACTGTGTCTACTGTCTGGAAGACATGTC 4140
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Db       4116 ATCTTCCCTGTCAGGCTCTTACAGCCACAGGCACTGTGTCTACTGTCTGGAAGACATGTC 4175

Qy       4141 CCCGTGGCTGTGGGGCCGCTGCTTCTGTTTAAATAAAAGTGGCCTGG 4187
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RESULT 13
US-10-087-192-1259
; Sequence 1259, Application US/10087192
; Publication No. US20020182586A1
; GENERAL INFORMATION:
; APPLICANT: Morris, David W.
; APPLICANT: Engelhard, Eric K.
; TITLE OF INVENTION: NOVEL COMPOSITIONS AND METHODS FOR
; TITLE OF INVENTION: CANCER
; FILE REFERENCE: 529452000122
; CURRENT APPLICATION NUMBER: US/10/087,192
; CURRENT FILING DATE: 2002-03-01
; PRIOR APPLICATION NUMBER: US 09/747,377
; PRIOR FILING DATE: 2000-12-22
; PRIOR APPLICATION NUMBER: US 09/798,586
; PRIOR FILING DATE: 2001-03-02
; NUMBER OF SEQ ID NOS: 2059
; SOFTWARE: FastSEQ for Windows Version 4.0
; SEQ ID NO 1259
; LENGTH: 4248
; TYPE: DNA
; ORGANISM: Homo sapiens
US-10-087-192-1259

Query Match 99.9%; Score 4182.2; DB 13; Length 4248;
Best Local Similarity 99.9%; Pred. No. 0;
Matches 4184; Conservative 0; Mismatches 3; Indels 0; Gaps 0;

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Db         62 GGCGCGAACCCGCAGCGCTTACCGCGCGGCGCCGCACCATGGAGCCCGCCGTGTCGCTGG 121

Qy         61 CCGTGTGCGCGCTGCTCTTCCTGCTGTGGGTGCGCCTGAAGGGGCTGGAGTTCGTGCTCA 120
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        122 CCGTGTGCGCGCTGCTCTTCCTGCTGTGGGTGCGCCTGAAGGGGCTGGAGTTCGTGCTCA 181

Qy        121 TCCACCAGCGCTGGGTGTTCGTGTGCCTCTTCCTCCTGCCGCTCTCGCTTATCTTCGATA 180
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        182 TCCACCAGCGCTGGGTGTTCGTGTGCCTCTTCCTCCTGCCGCTCTCGCTTATCTTCGATA 241

Qy        181 TCTACTACTACGTGCGCGCCTGGGTGGTGTTCAAGCTCAGCAGCGCTCCGCGCCTGCACG 240
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        242 TCTACTACTACGTGCGCGCCTGGGTGGTGTTCAAGCTCAGCAGCGCTCCGCGCCTGCACG 301

Qy        241 AGCAGCGCGTGCGGGACATCCAGAAGCAGGTGCGGGAATGGAAGGAGCAGGGTAGCAAGA 300
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        302 AGCAGCGCGTGCGGGACATCCAGAAGCAGGTGCGGGAATGGAAGGAGCAGGGTAGCAAGA 361
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Db        576 TCATGGGCACAGGCATCGAGTCATCATCCCACAAGTACGGCCTGTTCCAACACATCTGCA 635
Qy        601 CTGCTTACGAGCTGGTCCTGGCTGATGGCAGCTTTGTGCGATGCACTCCGTCCGAAAACT 660
Db        636 CTGCTTACGAGCTGGTCCTGGCTGATGGCAGCTTTGTGCGATGCACTCCGTCCGAAAACT 695
Qy        661 CAGACCTGTTCTATGCCGTACCCTGGTCCTGTGGGACGCTGGGTTTCCTGGTGGCCGCTG 720
Db        696 CAGACCTGTTCTATGCCGTACCCTGGTCCTGTGGGACGCTGGGTTTCCTGGTGGCCGCTG 755
Qy        721 AGATCCGCATCATCCCTGCCAAGAAGTACGTCAAGCTGCGTTTCGAGCCAGTGCGGGGCC 780
Db        756 AGATCCGCATCATCCCTGCCAAGAAGTACGTCAAGCTGCGTTTCGAGCCAGTGCGGGGCC 815
Qy        781 TGGAGGCTATCTGTGCCAAGTTCACCCACGAGTCCCAGCGGCAGGAGAACCACTTCGTGG 840
Db        816 TGGAGGCTATCTGTGCCAAGTTCACCCACGAGTCCCAGCGGCAGGAGAACCACTTCGTGG 875
Qy        841 AAGGGCTGCTCTACTCCCTGGATGAGGCTGTCATTATGACAGGGGTCATGACAGATGAGG 900
Db        876 AAGGGCTGCTCTACTCCCTGGATGAGGCTGTCATTATGACAGGGGTCATGACAGATGAGG 935
Qy        901 CAGAGCCCAGCAAGCTGAATAGCATTGGCAATTACTACAAGCCGTGGTTCTTTAAGCATG 960
Db        936 CAGAGCCCAGCAAGCTGAATAGCATTGGCAATTACTACAAGCCGTGGTTCTTTAAGCATG 995
Qy        961 TGGAGAACTATCTGAAGACAAACCGAGAGGGCCTGGAGTACATTCCCTTGAGACACTACT 1020
Db        996 TGGAGAACTATCTGAAGACAAACCGAGAGGGCCTGGAGTACATTCCCTTGAGACACTACT 1055
Qy       1021 ACCACCGCCACACGCGCAGCATCTTCTGGGAGCTCCAGGACATCATCCCCTTTGGCAACA 1080
Db       1056 ACCACCGCCACACGCGCAGCATCTTTTGGGAGCTCCAGGACATCATCCCCTTTGGCAACA 1115
Qy       1081 ACCCCATCTTCCGCTACCTCTTTGGCTGGATGGTGCCTCCCAAGATCTCCCTCCTGAAGC 1140
Db       1116 ACCCCATCTTCCGCTACCTCTTTGGCTGGATGGTGCCTCCCAAGATCTCCCTCCTGAAGC 1175
Qy       1141 TGACCCAGGGTGAGACCCTGCGCAAGCTGTACGAGCAGCACCACGTGGTGCAGGACATGC 1200
Db       1176 TGACCCAGGGTGAGACCCTGCGCAAGCTGTACGAGCAGCACCACGTGGTGCAGGACATGC 1235
Qy       1201 TGGTGCCCATGAAGTGCCTGCAGCAGGCCCTGCACACCTTCCAAAACGACATCCACGTCT 1260
Db       1236 TGGTGCCCATGAAGTGCCTGCAGCAGGCCCTGCACACCTTCCAAAACGACATCCACGTCT 1295
Qy       1261 ACCCCATCTGGCTGTGTCCGTTCATCCTGCCCAGCCAGCCAGGCCTAGTGCACCCCAAAG 1320
Db       1296 ACCCCATCTGGCTGTGTCCGTTCATCCTGCCCAGCCAGCCAGGCCTAGTGCACCCCAAAG 1355
Qy       1321 GAAATGAGGCAGAGCTCTACATCGACATTGGAGCATATGGGGAGCCGCGTGTGAAACACT 1380
Db       1356 GAAATGAGGCAGAGCTCTACATCGACATTGGAGCATATGGGGAGCCGCGTGTGAAACACT 1415
Qy       1381 TTGAAGCCAGGTCCTGCATGAGGCAGCTGGAGAAGTTTGTCCGCAGCGTGCATGGCTTCC 1440
Db       1416 TTGAAGCCAGGTCCTGCATGAGGCAGCTGGAGAAGTTTGTCCGCAGCGTGCATGGCTTCC 1475
Qy       1441 AGATGCTGTATGCCGACTGCTACATGAACCGGGAGGAGTTCTGGGAGATGTTTGATGGCT 1500
Db       1476 AGATGCTGTATGCCGACTGCTACATGAACCGGGAGGAGTTCTGGGAGATGTTTGATGGCT 1535
Qy       1501 CCTTGTACCACAAGCTGCGAGAGAAGCTGGGTTGCCAGGACGCCTTCCCCGAGGTGTACG 1560
Db       1536 CCTTGTACCACAAGCTGCGAGAGAAGCTGGGTTGCCAGGACGCCTTCCCCGAGGTGTACG 1595
Qy       1561 ACAAGATCTGCAAGGCCGCCAGGCACTGAGCTGGAGCCCGCCTGGAGAGACAGACACGTG 1620
Db       1596 ACAAGATCTGCAAGGCCGCCAGGCACTGAGCTGGAGCCCGCCTGGAGAGACAGACACGTG 1655
Qy       1621 TGAGTGGTCAGGCATCTTCCCTTCACTCAAGCTTGGCTGCTTTCCTAGATCCACACTTTC 1680
Db       1656 TGAGTGGTCAGGCATCTTCCCTTCACTCAAGCTTGGCTGCTTTCCTAGATCCACACTTTC 1715
Qy       1681 AAAGAGAAACCCCTCCAGAACTCCCACCCTGACAGCCCAACACCACCTTCCTCCTGGCTT 1740
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Qy       1741 CCAGGGGGCAGCCCAGTGGAATGGAAAGAATGTGGGATTTGGAGTCAGACAAGCCTGAGT 1800
Db       1776 CCAGGGGGCAGCCCAGTGGAATGGAAAGAATGTGGGATTTGGAGTCAGACAAGCCTGAGT 1835
Qy       1801 CCAGTTCCCCGTTTAGAACTCATTAGCTGTGTGACTCTGGGTGAGTCCCTTAACCCCTCT 1860
Db       1836 CCAGTTCCCCGTTTAGAACTCATTAGCTGTGTGACTCTGGGTGAGTCCCTTAACCCCTCT 1895
Qy       1861 GAGCCCGGGTCTCTTCATTAGTTGAAAGGGATAGTAATACCTACTTGCAGGTTGTTGTCA 1920
Db       1896 GAGCCCGGGTCTCTTCATTAGTTGAAAGGGATAGTAATACCTACTTGCAGGTTGTTGTCA 1955
Qy       1921 TCTGAGTTGAGCACTGGTCACATTGAAGGTGCTGGGTAAGTGGTAGCTCTTGTTGCTTCC 1980
Db       1956 TCTGAGTTGAGCACTGGTCACATTGAAGGTGCTGGGTAAGTGGTAGCTCTTGTTGCTTCC 2015
Qy       1981 CGTTCAGCGTCACATCTGCAGTGGAGCCTGAAAAGGCTCCACATTAGGTCACCTGTGCAC 2040
Db       2016 CGTTCAGCGTCACATCTGCAGTGGAGCCTGAAAAGGCTCCACATTAGGTCACCTGTGCAC 2075
Qy       2041 AGCCATGGCTGGAATGATGAAGGGGATACGCTGGAGTTGCCCTGCCATCGCCTCCATCAG 2100
Db       2076 AGCCATGGCTGGAATGATGAAGGGGATACGCTGGAGTTGCCCTGCCATCGCCTCCATCAG 2135
Qy       2101 CCAGACGAGGTCCTCACAGGAGAAGGACAGCTCTTCCCCACCCTGGGATCTCAGGAGGGC 2160
Db       2136 CCAGACGAGGTCCTCACAGGAGAAGGACAGCTCTTCCCCACCCTGGGATCTCAGGAGGGC 2195
Qy       2161 AGCCACGGAGTGGGGAGGCCCCAGATGCGCTGTGCCAAAGCCAGGTCCGAGGCCAAAGTT 2220
Db       2196 AGCCACGGAGTGGGGAGGCCCCAGATGCGCTGTGCCAAAGCCAGGTCCGAGGCCAAAGTT 2255
Qy       2221 CTCCCTGCCATCCTTGGTGCCGTCCTGCCCCTTCCTCCTTCATGCCTGGGCCTGCAGGCC 2280
Db       2256 CTCCCTGCCATCCTTGGTGCCGTCCTGCCCCTTCCTCCTTCATGCCTGGGCCTGCAGGCC 2315
Qy       2281 CACCCCAGCCACCACTGAGTCCACTCGGAGTGCCCTGTGTTCCTGGAGAAGGCATTCCAG 2340
Db       2316 CACCCCAGCCACCACTGAGTCCACTCGGAGTGCCCTGTGTTCCTGGAGAAGGCATTCCAG 2375
Qy       2341 GGTTGAATCTTGTCCCAGCCTCAGCCTGGGACACCTAGGTGGAGAGAGTGGTCTCCGCTC 2400
Db       2376 GGTTGAATCTTGTCCCAGCCTCAGCCTGGGACACCTAGGTGGAGAGAGTGGTCTCCGCTC 2435
Qy       2401 TGAATTGGATCCAGGGGACCTGGGCTCATTCTTCTTGGCTCACCAACCCTGCAGGCCTCA 2460
Db       2436 TGAATTGGATCCAGGGGACCTGGGCTCATTCTTCTTGGCTCACCAACCCTGCAGGCCTCA 2495
Qy       2461 TCTTTCCCAAAACCCACTTTGTCTTGGTGGGAGTGGGTCCGCGCTGCTCTGCAGCAGGGG 2520
Db       2496 TCTTTCCCAAAACCCACTTTGTCTTGGTGGGAGTGGGTCCGCGCTGCTCTGCAGCAGGGG 2555
Qy       2521 CTGGGGAGTGGACAGCATCAGGTGGGAAAGTGGAGTCCACCCTCATGTTTCTGTAGGATT 2580
Db       2556 CTGGGGAGTGGACAGCATCAGGTGGGAAAGTGGAGTCCACCCTCATGTTTCTGTAGGATT 2615
Qy       2581 CTCACCGTGGGGCTGGAAGAAAAGAGCATCGACTTGATTTCTCCAACCACTCATCCCTCT 2640
Db       2616 CTCACCGTGGGGCTGGAAGAAAAGAGCATCGACTTGATTTCTCCAACCACTCATCCCTCT 2675
Qy       2641 TTTTCTTTCTTCCACCACTCCCCACCCCAGCTGTAGTTAATTTCAGTGCCTTACAAATCC 2700
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Qy       2701 TAAGCTCAGAGAAAGTTCCATTTCCGTTCCAGAGGGAAGGGAACCTCCCTAGGTCCTTCC 2760
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Db       3062 CATCTGTCACATCGCTACAGACAAGACCTCCCTGGAACCCTTCTGATTGTCTTAGACACT 3121

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Qy       3121 GTGGACACTTGGGGATTATCTGTCATTCAAGATCCTTCCTTCAACCCCAAGGCCAGCTCC 3180

Db       3182 GTGGACACTTGGGGATTATCTGTCATTCAAGATCCTTCCTTCAACCCCAAGGCCAGCTCC 3241

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Qy       3421 ATCCTATATTTTACAAAGCATGTGAATTCTGGCATTAGCTCTCATAGGAGACCCATGTGC 3480

Db       3482 ATCCTATATTTTACAAAGCATGTGAATTCTGGCATTAGCTCTCATAGGAGACCCATGTGC 3541

Qy       3481 TTCCTTGCTCAGTGCAAAACTGATGATTCTACTTGCTGTAGATGAATGGTTAACACGAGC 3540

Db       3542 TTCCTTGCTCAGTGCAAAACTGATGATTCTACTTGCTGTAGATGAATGGTTAACACGAGC 3601

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Db       3602 TAGTTAAACAGTGCCATTGTTTTGCCAGTGAAGCCTCCAACCCTAAGCCACTGGGACGGT 3661

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Qy       3661 CCACAACCCGGGGCTTGGAAAGGAAGGTATTTTGGAATCACACCCTCCGGTTATGTTGCT 3720

Db       3722 CCACAACCCGGGGCTTGGAAAGGAAGGTATTTTGGAATCACACCCTCCGGTTATGTTGCT 3781

Qy       3721 CCAGTAAAATCTTGCCTGGAAAGAGGCAGTCTTCTTAGCATGGTGAGCTGAGTTCATGGC 3780

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Db       3842 TTTTTTTTGTAGCCAGTCCTGTCCCTGGCCATCCATGTGATGGTTTTGGATGGAGTTAAA 3901

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Qy       4141 CCCGTGGCTGTGGGGCCGCTGCTTCTGTTTAAATAAAAGTGGCCTGG 4187

Db       4202 CCCGTGGCTGTGGGGCCGCTGCTTCTGTTTAAATAAAAGTGGCCTGG 4248
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RESULT 12
US-10-176-847-1
; Sequence 1, Application US/10176847
; Publication No. US20030068636A1
; GENERAL INFORMATION:
; APPLICANT: Veiby, Petter Ole
; TITLE OF INVENTION: COMPOSITIONS, KITS, AND METHODS FOR
; TITLE OF INVENTION: IDENTIFICATION, ASSESSMENT, PREVENTION, AND THERAPY OF BREAST
; TITLE OF INVENTION: AND OVARIAN CANCER
; FILE REFERENCE: MRI-039
; CURRENT APPLICATION NUMBER: US/10/176,847
; CURRENT FILING DATE: 2002-06-21
; NUMBER OF SEQ ID NOS: 112
; SOFTWARE: FastSEQ for Windows Version 4.0
; SEQ ID NO 1
; LENGTH: 4275
; TYPE: DNA
; ORGANISM: Homo sapiens
; FEATURE:
; NAME/KEY: misc_feature
; LOCATION: 4241, 4243, 4244, 4246, 4247, 4250, 4253, 4254, 4255, 4259,
; LOCATION: 4260, 4261, 4262, 4263, 4266, 4270, 4271, 4272, 4273
; OTHER INFORMATION: n = A,T,C or G
US-10-176-847-1

Query Match 99.9%; Score 4183.8; DB 14; Length 4275;
Best Local Similarity 100.0%; Pred. No. 0;
Matches 4185; Conservative 0; Mismatches 2; Indels 0; Gaps 0;

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Qy        541 TCATGGGCACAGGCATCGAGTCATCATCCCACAAGTACGGCCTGTTCCAACACATCTGCA 600
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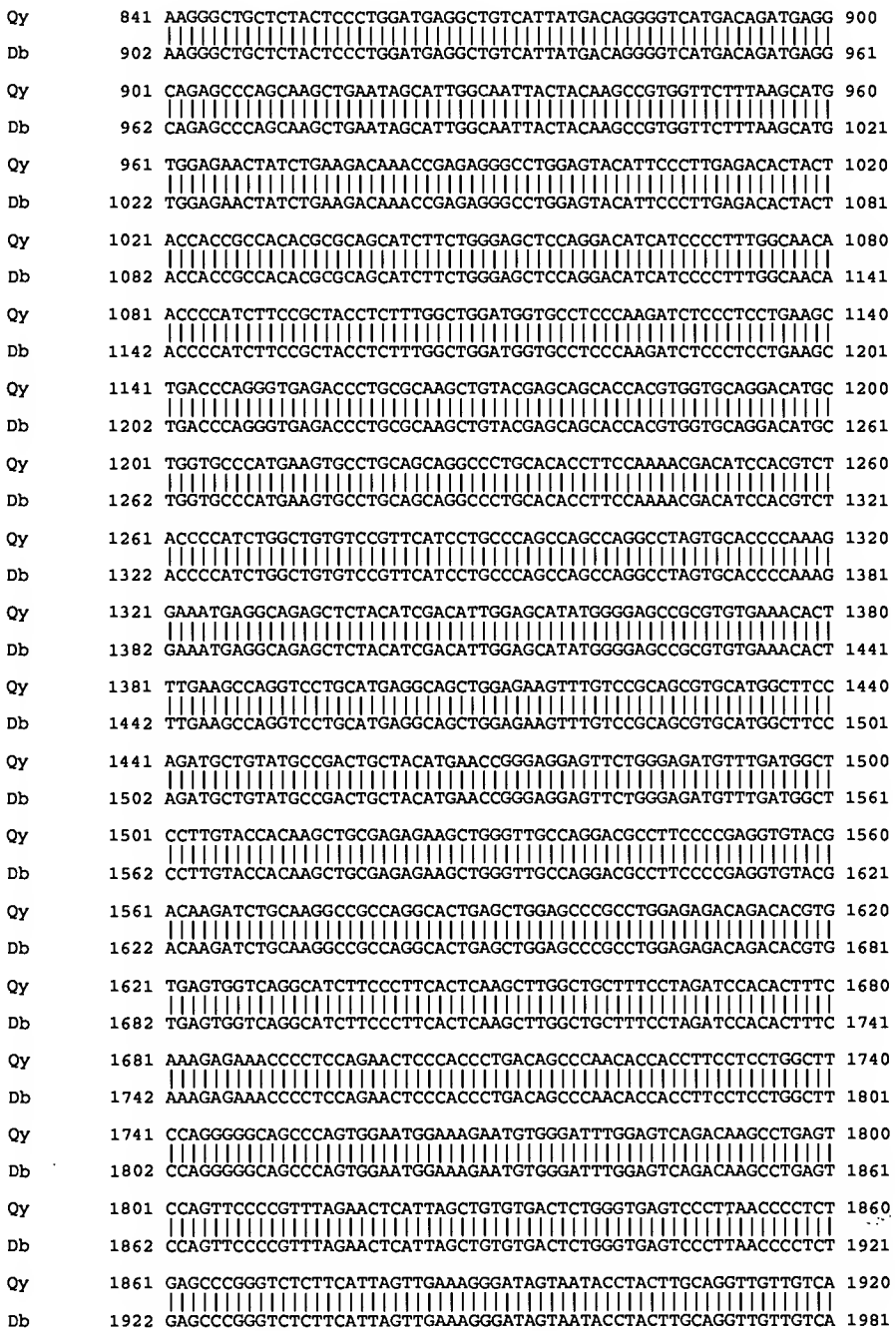
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Db 902 AAGGGCTGCTCTACTCCCTGGATGAGGCTGTCATTATGACAGGGGTCATGACAGATGAGG 961
Qy 901 CAGAGCCCAGCAAGCTGAATAGCATTGGCAATTACTACAAGCCGTGGTTCTTTAAGCATG 960
Db 962 CAGAGCCCAGCAAGCTGAATAGCATTGGCAATTACTACAAGCCGTGGTTCTTTAAGCATG 1021
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Db 1082 ACCACCGCCACACGCGCAGCATCTTCTGGGAGCTCCAGGACATCATCCCCTTTGGCAACA 1141
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Db 1142 ACCCCATCTTCCGCTACCTCTTTGGCTGGATGGTGCCTCCCAAGATCTCCCTCCTGAAGC 1201
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Db 1382 GAAATGAGGCAGAGCTCTACATCGACATTGGAGCATATGGGGAGCCGCGTGTGAAACACT 1441
Qy 1381 TTGAAGCCAGGTCCTGCATGAGGCAGCTGGAGAAGTTTGTCCGCAGCGTGCATGGCTTCC 1440
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Qy 1801 CCAGTTCCCCGTTTAGAACTCATTAGCTGTGTGACTCTGGGTGAGTCCCTTAACCCCTCT 1860
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Db 1922 GAGCCCGGGTCTCTTCATTAGTTGAAAGGGATAGTAATACCTACTTGCAGGTTGTTGTCA 1981
Qy 1921 TCTGAGTTGAGCACTGGTCACATTGAAGGTGCTGGGTAAGTGGTAGCTCTTGTTGCTTCC 1980
Db 1982 TCTGAGTTGAGCACTGGTCACATTGAAGGTGCTGGGTAAGTGGTAGCTCTTGTTGCTTCC 2041
Qy 1981 CGTTCAGCGTCACATCTGCAGTGGAGCCTGAAAAGGCTCCACATTAGGTCACCTGTGCAC 2040
Db 2042 CGTTCAGCGTCACATCTGCAGTGGAGCCTGAAAAGGCTCCACATTAGGTCACCTGTGCAC 2101
Qy 2041 AGCCATGGCTGGAATGATGAAGGGGATACGCTGGAGTTGCCCTGCCATCGCCTCCATCAG 2100
Db 2102 AGCCATGGCTGGAATGATGAAGGGGATACGCTGGAGTTGCCCTGCCATCGCCTCCATCAG 2161
Qy 2101 CCAGACGAGGTCCTCACAGGAGAAGGACAGCTCTTCCCCACCCTGGGATCTCAGGAGGGC 2160
Db 2162 CCAGACGAGGTCCTCACAGGAGAAGGACAGCTCTTCCCCACCCTGGGATCTCAGGAGGGC 2221
Qy 2161 AGCCACGGAGTGGGGAGGCCCCAGATGCGCTGTGCCAAAGCCAGGTCCGAGGCCAAAGTT 2220
Db 2222 AGCCACGGAGTGGGGAGGCCCCAGATGCGCTGTGCCAAAGCCAGGTCCGAGGCCAAAGTT 2281
Qy 2221 CTCCCTGCCATCCTTGGTGCCGTCCTGCCCCTTCCTCCTTCATGCCTGGGCCTGCAGGCC 2280
Db 2282 CTCCCTGCCATCCTTGGTGCCGTCCTGCCCCTTCCTCCTTCATGCCTGGGCCTGCAGGCC 2341
Qy 2281 CACCCCAGCCACCACTGAGTCCACTCGGAGTGCCCTGTGTTCCTGGAGAAGGCATTCCAG 2340
Db 2342 CACCCCAGCCACCACTGAGTCCACTCGGAGTGCCCTGTGTTCCTGGAGAAGGCATTCCAG 2401
Qy 2341 GGTTGAATCTTGTCCCAGCCTCAGCCTGGGACACCTAGGTGGAGAGAGTGGTCTCCGCTC 2400
Db 2402 GGTTGAATCTTGTCCCAGCCTCAGCCTGGGACACCTAGGTGGAGAGAGTGGTCTCCGCTC 2461
Qy 2401 TGAATTGGATCCAGGGGACCTGGGCTCATTCTTCTTGGCTCACCAACCCTGCAGGCCTCA 2460
Db 2462 TGAATTGGATCCAGGGGACCTGGGCTCATTCTTCTTGGCTCACCAACCCTGCAGGCCTCA 2521
Qy 2461 TCTTTCCCAAAACCCACTTTGTCTTGGTGGGAGTGGGTCCGCGCTGCTCTGCAGCAGGGG 2520
Db 2522 TCTTTCCCAAAACCCACTTTGTCTTGGTGGGAGTGGGTCCGCGCTGCTCTGCAGCAGGGG 2581
Qy 2521 CTGGGGAGTGGACAGCATCAGGTGGGAAAGTGGAGTCCACCCTCATGTTTCTGTAGGATT 2580
Db 2582 CTGGGGAGTGGACAGCATCAGGTGGGAAAGTGGAGTCCACCCTCATGTTTCTGTAGGATT 2641
Qy 2581 CTCACCGTGGGGCTGGAAGAAAAGAGCATCGACTTGATTTCTCCAACCACTCATCCCTCT 2640
Db 2642 CTCACCGTGGGGCTGGAAGAAAAGAGCATCGACTTGATTTCTCCAACCACTCATCCCTCT 2701
Qy 2641 TTTTCTTTCTTCCACCACTCCCCACCCCAGCTGTAGTTAATTTCAGTGCCTTACAAATCC 2700
Db 2702 TTTTCTTTCTTCCACCACTCCCCACCCCAGCTGTAGTTAATTTCAGTGCCTTACAAATCC 2761
Qy 2701 TAAGCTCAGAGAAAGTTCCATTTCCGTTCCAGAGGGAAGGGAACCTCCCTAGGTCCTTCC 2760
Db 2762 TAAGCTCAGAGAAAGTTCCATTTCCGTTCCAGAGGGAAGGGAACCTCCCTAGGTCCTTCC 2821
Qy 2761 CTGGCTTGTTATAACGCAAAGCTTGGTTGTTTATGCAACTCTATCTTAAGAACTGCCCAG 2820
Db 2822 CTGGCTTGTTATAACGCAAAGCTTGGTTGTTTATGCAACTCTATCTTAAGAACTGCCCAG 2881
Qy 2821 CCTCAGCTGAAAACCCGAATCTGAGAAGGAATTGCGTCATGTAAGGGAAGCTGGAATTAA 2880
Db 2882 CCTCAGCTGAAAACCCGAATCTGAGAAGGAATTGCGTCATGTAAGGGAAGCTGGAATTAA 2941
Qy 2881 GGGAGCTGAGCCAGTCATGGTTGTGGCGTGTGAGTCAGGAGACCTAGGTTTCAGCCCCTC 2940
Db 2942 GGGAGCTGAGCCAGTCATGGTTGTGGCGTGTGAGTCAGGAGACCTAGGTTTCAGCCCCTC 3001
Qy 2941 TCTACTGTCAGCGAGCTGTGCAACGTGGGCAAGTCATTGTCCTCTGAGCTGCAGTTTCCT 3000
Db 3002 TCTACTGTCAGCGAGCTGTGCAACGTGGGCAAGTCATTGTCCTCTGAGCTGCAGTTTCCT 3061
Qy 3001 CATCTGTCACATCGCTACAGACAAGACCTCCCTGGAACCCTTCTGATTGTCTTAGACACT 3060

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Qy       3421 ATCCTATATTTTACAAAGCATGTGAATTCTGGCATTAGCTCTCATAGGAGACCCATGTGC 3480
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Db       3482 ATCCTATATTTTACAAAGCATGTGAATTCTGGCATTAGCTCTCATAGGAGACCCATGTGC 3541

Qy       3481 TTCCTTGCTCAGTGCAAAACTGATGATTCTACTTGCTGTAGATGAATGGTTAACACGAGC 3540
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       3542 TTCCTTGCTCAGTGCAAAACTGATGATTCTACTTGCTGTAGATGAATGGTTAACACGAGC 3601

Qy       3541 TAGTTAAACAGTGCCATTGTTTTGCCAGTGAAGCCTCCAACCCTAAGCCACTGGGACGGT 3600
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Db       3602 TAGTTAAACAGTGCCATTGTTTTGCCAGTGAAGCCTCCAACCCTAAGCCACTGGGACGGT 3661

Qy       3601 GGCCAGAGATGCCAGCAGCCTCTGTCGCCCTTAGTCATATAACCAAAATCCAGACCTTAT 3660
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       3662 GGCCAGAGATGCCAGCAGCCTCTGTCGCCCTTAGTCATATAACCAAAATCCAGACCTTAT 3721

Qy       3661 CCACAACCCGGGGCTTGGAAAGGAAGGTATTTTGGAATCACACCCTCCGGTTATGTTGCT 3720
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       3722 CCACAACCCGGGGCTTGGAAAGGAAGGTATTTTGGAATCACACCCTCCGGTTATGTTGCT 3781

Qy       3721 CCAGTAAAATCTTGCCTGGAAAGAGGCAGTCTTCTTAGCATGGTGAGCTGAGTTCATGGC 3780
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       3782 CCAGTAAAATCTTGCCTGGAAAGAGGCAGTCTTCTTAGCATGGTGAGCTGAGTTCATGGC 3841

Qy       3781 TTTTTTTTGTAGCCAGTCCTGTCCCTGGCCATCCATGTGATGGTTTTGGATGGAGTTAAA 3840
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       3842 TTTTTTTTGTAGCCAGTCCTGTCCCTGGCCATCCATGTGATGGTTTTGGATGGAGTTAAA 3901

Qy       3841 CTTGATGCCAGTGGGCAGTGCATGTGGAAAGTATCAGAGTAAGCCTCTCCCCTCCAGAGC 3900
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       3902 CTTGATGCCAGTGGGCAGTGCATGTGGAAAGTATCAGAGTAAGCCTCTCCCCTCCAGAGC 3961

Qy       3901 CCTGAGTTTCTTGGCTGCATGAAGGTTTTCTTTAGAATCAGAATTGTAGCCAGTTTCTTT 3960
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Db       3962 CCTGAGTTTCTTGGCTGCATGAAGGTTTTCTTTAGAATCAGAATTGTAGCCAGTTTCTTT 4021

Qy       3961 GGCCAGAAGGATGAAATACTTGGATATTACTGAAAGGGAGGGGTGGAGATGGGTGTGGCAG 4020
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       4022 GGCCAGAAGGATGAAATACTTGGATATTACTGAAAGGGAGGGGTGGAGATGGGTGTGGCAG 4081

Qy       4021 TGTATGGTGTGTGATTTTTATTTTCTTCTTTGGTCATGGGGGCCAAGGAGAAAGGCATGA 4080
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       4082 TGTATGGTGTGTGATTTTTATTTTCTTCTTTGGTCATGGGGGCCAAGGAGAAAGGCATGA 4141

Qy       4081 ATCTTCCCTGTCAGGCTCTTACAGCCACAGGCACTGTGTCTACTGTCTGGAAGACATGTC 4140
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       4142 ATCTTCCCTGTCAGGCTCTTACAGCCACAGGCACTGTGTCTACTGTCTGGAAGACATGTC 4201

Qy       4141 CCCGTGGCTGTGGGGCCGCTGCTTCTGTTTAAATAAAAGTGGCCTGG 4187
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Db       4202 CCCGTGGCTGTGGGGCCGCTGCTTCTGTTTAAATAAAAGTGGCCTGG 4248
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RESULT 11
US-10-934-998-32
; Sequence 32, Application US/10934998
; Publication No. US20050153917A1
; GENERAL INFORMATION:
;  APPLICANT: AL-MAHMOOD, SALMAN
;  APPLICANT:  COLIN, SYLVIE
;  APPLICANT:  SCHNEIDER, CHRISTOPHE
;  TITLE OF INVENTION: GENES INVOLVED IN REGULATING ANGIOGENESIS, PHARMACEUTICAL
;  TITLE OF INVENTION:  PREPARATIONS CONTAINING SAME AND APPLICATIONS THEREOF
;  FILE REFERENCE: BMA-04-1206
;  CURRENT APPLICATION NUMBER: US/10/934,998
;  CURRENT FILING DATE:  2004-09-03
;  PRIOR APPLICATION NUMBER: PCT/FR03/00695
;  PRIOR FILING DATE: 2003-03-04
;  PRIOR APPLICATION NUMBER: FR02/02717
;  PRIOR FILING DATE: 2002-03-04
;  PRIOR APPLICATION NUMBER: FR02/04546
;  PRIOR FILING DATE: 2002-04-11
;  NUMBER OF SEQ ID NOS: 301
;  SOFTWARE: PatentIn version 3.2
; SEQ ID NO 32
;   LENGTH: 4248
;   TYPE: DNA
;   ORGANISM: Homo sapiens
;   FEATURE:
;   OTHER INFORMATION: GS-N31
;  PUBLICATION INFORMATION:
;   DATABASE ACCESSION NUMBER: GENBANK/AF261758
;   DATABASE ENTRY DATE: 2000-10-10
;   RELEVANT RESIDUES: (1)..(4248)
US-10-934-998-32

  Query Match             100.0%;  Score 4187;  DB 22;  Length 4248;
  Best Local Similarity   100.0%;  Pred. No. 0;
  Matches 4187;  Conservative    0;  Mismatches    0;  Indels    0;  Gaps    0;

Qy          1 GGCGCGAACCCGCAGCGCTTACCGCGCGGCGCCGCACCATGGAGCCCGCCGTGTCGCTGG 60
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Db         62 GGCGCGAACCCGCAGCGCTTACCGCGCGGCGCCGCACCATGGAGCCCGCCGTGTCGCTGG 121

Qy         61 CCGTGTGCGCGCTGCTCTTCCTGCTGTGGGTGCGCCTGAAGGGGCTGGAGTTCGTGCTCA 120
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        122 CCGTGTGCGCGCTGCTCTTCCTGCTGTGGGTGCGCCTGAAGGGGCTGGAGTTCGTGCTCA 181

Qy        121 TCCACCAGCGCTGGGTGTTCGTGTGCCTCTTCCTCCTGCCGCTCTCGCTTATCTTCGATA 180
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        182 TCCACCAGCGCTGGGTGTTCGTGTGCCTCTTCCTCCTGCCGCTCTCGCTTATCTTCGATA 241

Qy        181 TCTACTACTACGTGCGCGCCTGGGTGGTGTTCAAGCTCAGCAGCGCTCCGCGCCTGCACG 240
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        242 TCTACTACTACGTGCGCGCCTGGGTGGTGTTCAAGCTCAGCAGCGCTCCGCGCCTGCACG 301

Qy        241 AGCAGCGCGTGCGGGACATCCAGAAGCAGGTGCGGGAATGGAAGGAGCAGGGTAGCAAGA 300
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        302 AGCAGCGCGTGCGGGACATCCAGAAGCAGGTGCGGGAATGGAAGGAGCAGGGTAGCAAGA 361

Qy        301 CCTTCATGTGCACGGGGCGCCCTGGCTGGCTCACTGTCTCACTACGTGTCGGGAAGTACA 360
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        362 CCTTCATGTGCACGGGGCGCCCTGGCTGGCTCACTGTCTCACTACGTGTCGGGAAGTACA 421

Qy        361 AGAAGACACACAAAAACATCATGATCAACCTGATGGACATTCTGGAAGTGGACACCAAGA 420
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        422 AGAAGACACACAAAAACATCATGATCAACCTGATGGACATTCTGGAAGTGGACACCAAGA 481

Qy        421 AACAGATTGTCCGTGTGGAGCCCTTGGTGACCATGGGCCAGGTGACTGCCCTGCTGACCT 480
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        482 AACAGATTGTCCGTGTGGAGCCCTTGGTGACCATGGGCCAGGTGACTGCCCTGCTGACCT 541

Qy        481 CCATTGGCTGGACTCTCCCCGTGTTGCCTGAGCTTGATGACCTCACAGTGGGGGGCTTGA 540
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        542 CCATTGGCTGGACTCTCCCCGTGTTGCCTGAGCTTGATGACCTCACAGTGGGGGGCTTGA 601

Qy        541 TCATGGGCACAGGCATCGAGTCATCATCCCACAAGTACGGCCTGTTCCAACACATCTGCA 600
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        602 TCATGGGCACAGGCATCGAGTCATCATCCCACAAGTACGGCCTGTTCCAACACATCTGCA 661

Qy        601 CTGCTTACGAGCTGGTCCTGGCTGATGGCAGCTTTGTGCGATGCACTCCGTCCGAAAACT 660
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        662 CTGCTTACGAGCTGGTCCTGGCTGATGGCAGCTTTGTGCGATGCACTCCGTCCGAAAACT 721

Qy        661 CAGACCTGTTCTATGCCGTACCCTGGTCCTGTGGGACGCTGGGTTTCCTGGTGGCCGCTG 720
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Db        722 CAGACCTGTTCTATGCCGTACCCTGGTCCTGTGGGACGCTGGGTTTCCTGGTGGCCGCTG 781

Qy        721 AGATCCGCATCATCCCTGCCAAGAAGTACGTCAAGCTGCGTTTCGAGCCAGTGCGGGGCC 780
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        782 AGATCCGCATCATCCCTGCCAAGAAGTACGTCAAGCTGCGTTTCGAGCCAGTGCGGGGCC 841

Qy        781 TGGAGGCTATCTGTGCCAAGTTCACCCACGAGTCCCAGCGGCAGGAGAACCACTTCGTGG 840
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        842 TGGAGGCTATCTGTGCCAAGTTCACCCACGAGTCCCAGCGGCAGGAGAACCACTTCGTGG 901
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Db      1262 TGGTGCCCATGAAGTGCCTGCAGCAGGCCCTGCACACCTTCCAAAACGACATCCACGTCT 1321
Qy      1261 ACCCCATCTGGCTGTGTCCGTTCATCCTGCCCAGCCAGCCAGGCCTAGTGCACCCCAAAG 1320
Db      1322 ACCCCATCTGGCTGTGTCCGTTCATCCTGCCCAGCCAGCCAGGCCTAGTGCACCCCAAAG 1381
Qy      1321 GAAATGAGGCAGAGCTCTACATCGACATTGGAGCATATGGGGAGCCGCGTGTGAAACACT 1380
Db      1382 GAAATGAGGCAGAGCTCTACATCGACATTGGAGCATATGGGGAGCCGCGTGTGAAACACT 1441
Qy      1381 TTGAAGCCAGGTCCTGCATGAGGCAGCTGGAGAAGTTTGTCCGCAGCGTGCATGGCTTCC 1440
Db      1442 TTGAAGCCAGGTCCTGCATGAGGCAGCTGGAGAAGTTTGTCCGCAGCGTGCATGGCTTCC 1501
Qy      1441 AGATGCTGTATGCCGACTGCTACATGAACCGGGAGGAGTTCTGGGAGATGTTTGATGGCT 1500
Db      1502 AGATGCTGTATGCCGACTGCTACATGAACCGGGAGGAGTTCTGGGAGATGTTTGATGGCT 1561
Qy      1501 CCTTGTACCACAAGCTGCGAGAGAAGCTGGGTTGCCAGGACGCCTTCCCCGAGGTGTACG 1560
Db      1562 CCTTGTACCACAAGCTGCGAGAGAAGCTGGGTTGCCAGGACGCCTTCCCCGAGGTGTACG 1621
Qy      1561 ACAAGATCTGCAAGGCCGCCAGGCACTGAGCTGGAGCCCGCCTGGAGAGACAGACACGTG 1620
Db      1622 ACAAGATCTGCAAGGCCGCCAGGCACTGAGCTGGAGCCCGCCTGGAGAGACAGACACGTG 1681
Qy      1621 TGAGTGGTCAGGCATCTTCCCTTCACTCAAGCTTGGCTGCTTTCCTAGATCCACACTTTC 1680
Db      1682 TGAGTGGTCAGGCATCTTCCCTTCACTCAAGCTTGGCTGCTTTCCTAGATCCACACTTTC 1741
Qy      1681 AAAGAGAAACCCCTCCAGAACTCCCACCCTGACAGCCCAACACCACCTTCCTCCTGGCTT 1740
Db      1742 AAAGAGAAACCCCTCCAGAACTCCCACCCTGACAGCCCAACACCACCTTCCTCCTGGCTT 1801
Qy      1741 CCAGGGGGCAGCCCAGTGGAATGGAAAGAATGTGGGATTTGGAGTCAGACAAGCCTGAGT 1800
Db      1802 CCAGGGGGCAGCCCAGTGGAATGGAAAGAATGTGGGATTTGGAGTCAGACAAGCCTGAGT 1861
Qy      1801 CCAGTTCCCCGTTTAGAACTCATTAGCTGTGTGACTCTGGGTGAGTCCCTTAACCCCTCT 1860
Db      1862 CCAGTTCCCCGTTTAGAACTCATTAGCTGTGTGACTCTGGGTGAGTCCCTTAACCCCTCT 1921
Qy      1861 GAGCCCGGGTCTCTTCATTAGTTGAAAGGGATAGTAATACCTACTTGCAGGTTGTTGTCA 1920
Db      1922 GAGCCCGGGTCTCTTCATTAGTTGAAAGGGATAGTAATACCTACTTGCAGGTTGTTGTCA 1981
Qy      1921 TCTGAGTTGAGCACTGGTCACATTGAAGGTGCTGGGTAAGTGGTAGCTCTTGTTGCTTCC 1980
Db      1982 TCTGAGTTGAGCACTGGTCACATTGAAGGTGCTGGGTAAGTGGTAGCTCTTGTTGCTTCC 2041
Qy      1981 CGTTCAGCGTCACATCTGCAGTGGAGCCTGAAAAGGCTCCACATTAGGTCACCTGTGCAC 2040
Db      2042 CGTTCAGCGTCACATCTGCAGTGGAGCCTGAAAAGGCTCCACATTAGGTCACCTGTGCAC 2101
Qy      2041 AGCCATGGCTGGAATGATGAAGGGGATACGCTGGAGTTGCCCTGCCATCGCCTCCATCAG 2100
Db      2102 AGCCATGGCTGGAATGATGAAGGGGATACGCTGGAGTTGCCCTGCCATCGCCTCCATCAG 2161
Qy      2101 CCAGACGAGGTCCTCACAGGAGAAGGACAGCTCTTCCCCACCCTGGGATCTCAGGAGGGC 2160
Db      2162 CCAGACGAGGTCCTCACAGGAGAAGGACAGCTCTTCCCCACCCTGGGATCTCAGGAGGGC 2221
Qy      2161 AGCCACGGAGTGGGGAGGCCCCAGATGCGCTGTGCCAAAGCCAGGTCCGAGGCCAAAGTT 2220
Db      2222 AGCCACGGAGTGGGGAGGCCCCAGATGCGCTGTGCCAAAGCCAGGTCCGAGGCCAAAGTT 2281
Qy      2221 CTCCCTGCCATCCTTGGTGCCGTCCTGCCCCTTCCTCCTTCATGCCTGGGCCTGCAGGCC 2280
Db      2282 CTCCCTGCCATCCTTGGTGCCGTCCTGCCCCTTCCTCCTTCATGCCTGGGCCTGCAGGCC 2341
Qy      2281 CACCCCAGCCACCACTGAGTCCACTCGGAGTGCCCTGTGTTCCTGGAGAAGGCATTCCAG 2340
Db      2342 CACCCCAGCCACCACTGAGTCCACTCGGAGTGCCCTGTGTTCCTGGAGAAGGCATTCCAG 2401
Qy      2341 GGTTGAATCTTGTCCCAGCCTCAGCCTGGGACACCTAGGTGGAGAGAGTGGTCTCCGCTC 2400
Db      2402 GGTTGAATCTTGTCCCAGCCTCAGCCTGGGACACCTAGGTGGAGAGAGTGGTCTCCGCTC 2461
Qy      2401 TGAATTGGATCCAGGGGACCTGGGCTCATTCTTCTTGGCTCACCAACCCTGCAGGCCTCA 2460
Db      2462 TGAATTGGATCCAGGGGACCTGGGCTCATTCTTCTTGGCTCACCAACCCTGCAGGCCTCA 2521
Qy      2461 TCTTTCCCAAAACCCACTTTGTCTTGGTGGGAGTGGGTCCGCGCTGCTCTGCAGCAGGGG 2520
Db      2522 TCTTTCCCAAAACCCACTTTGTCTTGGTGGGAGTGGGTCCGCGCTGCTCTGCAGCAGGGG 2581
Qy      2521 CTGGGGAGTGGACAGCATCAGGTGGGAAAGTGGAGTCCACCCTCATGTTTCTGTAGGATT 2580
Db      2582 CTGGGGAGTGGACAGCATCAGGTGGGAAAGTGGAGTCCACCCTCATGTTTCTGTAGGATT 2641
Qy      2581 CTCACCGTGGGGCTGGAAGAAAAGAGCATCGACTTGATTTCTCCAACCACTCATCCCTCT 2640
Db      2642 CTCACCGTGGGGCTGGAAGAAAAGAGCATCGACTTGATTTCTCCAACCACTCATCCCTCT 2701
Qy      2641 TTTTCTTTCTTCCACCACTCCCCACCCCAGCTGTAGTTAATTTCAGTGCCTTACAAATCC 2700
Db      2702 TTTTCTTTCTTCCACCACTCCCCACCCCAGCTGTAGTTAATTTCAGTGCCTTACAAATCC 2761
Qy      2701 TAAGCTCAGAGAAAGTTCCATTTCCGTTCCAGAGGGAAGGGAACCTCCCTAGGTCCTTCC 2760
Db      2762 TAAGCTCAGAGAAAGTTCCATTTCCGTTCCAGAGGGAAGGGAACCTCCCTAGGTCCTTCC 2821
Qy      2761 CTGGCTTGTTATAACGCAAAGCTTGGTTGTTTATGCAACTCTATCTTAAGAACTGCCCAG 2820
Db      2822 CTGGCTTGTTATAACGCAAAGCTTGGTTGTTTATGCAACTCTATCTTAAGAACTGCCCAG 2881
Qy      2821 CCTCAGCTGAAAACCCGAATCTGAGAAGGAATTGCGTCATGTAAGGGAAGCTGGAATTAA 2880
Db      2882 CCTCAGCTGAAAACCCGAATCTGAGAAGGAATTGCGTCATGTAAGGGAAGCTGGAATTAA 2941
Qy      2881 GGGAGCTGAGCCAGTCATGGTTGTGGCGTGTGAGTCAGGAGACCTAGGTTTCAGCCCCTC 2940
Db      2942 GGGAGCTGAGCCAGTCATGGTTGTGGCGTGTGAGTCAGGAGACCTAGGTTTCAGCCCCTC 3001
Qy      2941 TCTACTGTCAGCGAGCTGTGCAACGTGGGCAAGTCATTGTCCTCTGAGCTGCAGTTTCCT 3000
Db      3002 TCTACTGTCAGCGAGCTGTGCAACGTGGGCAAGTCATTGTCCTCTGAGCTGCAGTTTCCT 3061
Qy      3001 CATCTGTCACATCGCTACAGACAAGACCTCCCTGGAACCCTTCTGATTGTCTTAGACACT 3060
Db      3062 CATCTGTCACATCGCTACAGACAAGACCTCCCTGGAACCCTTCTGATTGTCTTAGACACT 3121
Qy      3061 GTGGTTGCAAAACCCACGGAAAGCCTCATTTGTGTGGAAAGTCAGAGGAAAAATGATCCA 3120
Db      3122 GTGGTTGCAAAACCCACGGAAAGCCTCATTTGTGTGGAAAGTCAGAGGAAAAATGATCCA 3181
Qy      3121 GTGGACACTTGGGGATTATCTGTCATTCAAGATCCTTCCTTCAACCCCAAGGCCAGCTCC 3180
Db      3182 GTGGACACTTGGGGATTATCTGTCATTCAAGATCCTTCCTTCAACCCCAAGGCCAGCTCC 3241
Qy      3181 CATCTCATTTCCAGAAAGGCTCATACCTGGCTTGCAGGGAAGCATCTGTCTTGTCATTCC 3240
Db      3242 CATCTCATTTCCAGAAAGGCTCATACCTGGCTTGCAGGGAAGCATCTGTCTTGTCATTCC 3301
Qy      3241 AGGTGCCAGAATCCTCTCAGAGTCATTGAAGGGTGTTCACCCATCCCACCCAAGGCTTGG 3300
Db      3302 AGGTGCCAGAATCCTCTCAGAGTCATTGAAGGGTGTTCACCCATCCCACCCAAGGCTTGG 3361
Qy      3301 CACACTGCCAGTGTCTTAGCAGGGTCTTGTGAGGGCTGGGGGCATCCAGGCACTCAGAAG 3360
Db      3362 CACACTGCCAGTGTCTTAGCAGGGTCTTGTGAGGGCTGGGGGCATCCAGGCACTCAGAAG 3421
Qy      3361 GCAAAGGAACCACCCTACCCATTTGGCCTCTGGAGGGGGCAGAAGAAAGAAAGAAACCTC 3420
Db      3422 GCAAAGGAACCACCCTACCCATTTGGCCTCTGGAGGGGGCAGAAGAAAGAAAGAAACCTC 3481
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Qy       3601 GGCCAGAGATGCCAGCAGCCTCTGTCGCCCTTAGTCATATAACCAAAATCCAGACCTTAT 3660
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Db       3601 GGCCAGAGATGCCAGCAGCCTCTGTCGCCCTTAGTCATATAACCAAAATCCAGACCTTAT 3660

Qy       3661 CCACAACCCGGGGCTTGGAAAGGAAGGTATTTTGGAATCACACCCTCCGGTTATGTTGCT 3720
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Db       3661 CCACAACCCGGGGCTTGGAAAGGAAGGTATTTTGGAATCACACCCTCCGGTTATGTTGCT 3720

Qy       3721 CCAGTAAAATCTTGCCTGGAAAGAGGCAGTCTTCTTAGCATGGTGAGCTGAGTTCATGGC 3780
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Db       3721 CCAGTAAAATCTTGCCTGGAAAGAGGCAGTCTTCTTAGCATGGTGAGCTGAGTTCATGGC 3780

Qy       3781 TTTTTTTTGTAGCCAGTCCTGTCCCTGGCCATCCATGTGATGGTTTTGGATGGAGTTAAA 3840
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       3781 TTTTTTTTGTAGCCAGTCCTGTCCCTGGCCATCCATGTGATGGTTTTGGATGGAGTTAAA 3840

Qy       3841 CTTGATGCCAGTGGGCAGTGCATGTGGAAAGTATCAGAGTAAGCCTCTCCCCTCCAGAGC 3900
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Db       3841 CTTGATGCCAGTGGGCAGTGCATGTGGAAAGTATCAGAGTAAGCCTCTCCCCTCCAGAGC 3900

Qy       3901 CCTGAGTTTCTTGGCTGCATGAAGGTTTTCTTTAGAATCAGAATTGTAGCCAGTTTCTTT 3960
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       3901 CCTGAGTTTCTTGGCTGCATGAAGGTTTTCTTTAGAATCAGAATTGTAGCCAGTTTCTTT 3960

Qy       3961 GGCCAGAAGGATGAATACTTGGATATTACTGAAAGGGAGGGGTGGAGATGGGTGTGGCAG 4020
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       3961 GGCCAGAAGGATGAATACTTGGATATTACTGAAAGGGAGGGGTGGAGATGGGTGTGGCAG 4020

Qy       4021 TGTATGGTGTGTGATTTTTATTTTCTTCTTTGGTCATGGGGGCCAAGGAGAAAGGCATGA 4080
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       4021 TGTATGGTGTGTGATTTTTATTTTCTTCTTTGGTCATGGGGGCCAAGGAGAAAGGCATGA 4080

Qy       4081 ATCTTCCCTGTCAGGCTCTTACAGCCACAGGCACTGTGTCTACTGTCTGGAAGACATGTC 4140
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       4081 ATCTTCCCTGTCAGGCTCTTACAGCCACAGGCACTGTGTCTACTGTCTGGAAGACATGTC 4140

Qy       4141 CCCGTGGCTGTGGGGCCGCTGCTTCTGTTTAAATAAAAGTGGCCTGG 4187
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Db       4141 CCCGTGGCTGTGGGGCCGCTGCTTCTGTTTAAATAAAAGTGGCCTGG 4187


RESULT 10
US-10-956-157-1950
; Sequence 1950, Application US/10956157
; Publication No. US20050118625A1
; GENERAL INFORMATION:
;  APPLICANT: Wyeth
;  APPLICANT:  Mounts, William
;  TITLE OF INVENTION: NUCLEIC ACID ARRAYS FOR DETECTING GENE EXPRESSION ASSOCIATED WITH
;  TITLE OF INVENTION:  HUMAN OSTEOARTHRITIS AND HUMAN PROTEASES
;  FILE REFERENCE: 031896-043000 (AM 101081)
;  CURRENT APPLICATION NUMBER: US/10/956,157
;  CURRENT FILING DATE:  2004-10-04
;  NUMBER OF SEQ ID NOS: 319805
;  SOFTWARE: PatentIn version 3.2
; SEQ ID NO 1950
;   LENGTH: 4248
;   TYPE: DNA
;   ORGANISM: Homo sapiens
US-10-956-157-1950

  Query Match            100.0%;  Score 4187;  DB 21;  Length 4248;
  Best Local Similarity  100.0%;  Pred. No. 0;
  Matches 4187;  Conservative    0;  Mismatches    0;  Indels    0;  Gaps    0;

Qy          1 GGCGCGAACCCGCAGCGCTTACCGCGCGGCGCCGCACCATGGAGCCCGCCGTGTCGCTGG 60
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db         62 GGCGCGAACCCGCAGCGCTTACCGCGCGGCGCCGCACCATGGAGCCCGCCGTGTCGCTGG 121

Qy         61 CCGTGTGCGCGCTGCTCTTCCTGCTGTGGGTGCGCCTGAAGGGGCTGGAGTTCGTGCTCA 120
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        122 CCGTGTGCGCGCTGCTCTTCCTGCTGTGGGTGCGCCTGAAGGGGCTGGAGTTCGTGCTCA 181

Qy        121 TCCACCAGCGCTGGGTGTTCGTGTGCCTCTTCCTCCTGCCGCTCTCGCTTATCTTCGATA 180
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Db        182 TCCACCAGCGCTGGGTGTTCGTGTGCCTCTTCCTCCTGCCGCTCTCGCTTATCTTCGATA 241

Qy        181 TCTACTACTACGTGCGCGCCTGGGTGGTGTTCAAGCTCAGCAGCGCTCCGCGCCTGCACG 240
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        242 TCTACTACTACGTGCGCGCCTGGGTGGTGTTCAAGCTCAGCAGCGCTCCGCGCCTGCACG 301

Qy        241 AGCAGCGCGTGCGGGACATCCAGAAGCAGGTGCGGGAATGGAAGGAGCAGGGTAGCAAGA 300
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        302 AGCAGCGCGTGCGGGACATCCAGAAGCAGGTGCGGGAATGGAAGGAGCAGGGTAGCAAGA 361

Qy        301 CCTTCATGTGCACGGGGCGCCCTGGCTGGCTCACTGTCTCACTACGTGTCGGGAAGTACA 360
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Db        362 CCTTCATGTGCACGGGGCGCCCTGGCTGGCTCACTGTCTCACTACGTGTCGGGAAGTACA 421

Qy        361 AGAAGACACACAAAAACATCATGATCAACCTGATGGACATTCTGGAAGTGGACACCAAGA 420
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        422 AGAAGACACACAAAAACATCATGATCAACCTGATGGACATTCTGGAAGTGGACACCAAGA 481

Qy        421 AACAGATTGTCCGTGTGGAGCCCTTGGTGACCATGGGCCAGGTGACTGCCCTGCTGACCT 480
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        482 AACAGATTGTCCGTGTGGAGCCCTTGGTGACCATGGGCCAGGTGACTGCCCTGCTGACCT 541

Qy        481 CCATTGGCTGGACTCTCCCCGTGTTGCCTGAGCTTGATGACCTCACAGTGGGGGGCTTGA 540
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        542 CCATTGGCTGGACTCTCCCCGTGTTGCCTGAGCTTGATGACCTCACAGTGGGGGGCTTGA 601

Qy        541 TCATGGGCACAGGCATCGAGTCATCATCCCACAAGTACGGCCTGTTCCAACACATCTGCA 600
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Db        602 TCATGGGCACAGGCATCGAGTCATCATCCCACAAGTACGGCCTGTTCCAACACATCTGCA 661

Qy        601 CTGCTTACGAGCTGGTCCTGGCTGATGGCAGCTTTGTGCGATGCACTCCGTCCGAAAACT 660
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Db        662 CTGCTTACGAGCTGGTCCTGGCTGATGGCAGCTTTGTGCGATGCACTCCGTCCGAAAACT 721

Qy        661 CAGACCTGTTCTATGCCGTACCCTGGTCCTGTGGGACGCTGGGTTTCCTGGTGGCCGCTG 720
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Db        722 CAGACCTGTTCTATGCCGTACCCTGGTCCTGTGGGACGCTGGGTTTCCTGGTGGCCGCTG 781

Qy        721 AGATCCGCATCATCCCTGCCAAGAAGTACGTCAAGCTGCGTTTCGAGCCAGTGCGGGGCC 780
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Qy        781 TGGAGGCTATCTGTGCCAAGTTCACCCACGAGTCCCAGCGGCAGGAGAACCACTTCGTGG 840
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Qy        841 AAGGGCTGCTCTACTCCCTGGATGAGGCTGTCATTATGACAGGGGTCATGACAGATGAGG 900
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Db        902 AAGGGCTGCTCTACTCCCTGGATGAGGCTGTCATTATGACAGGGGTCATGACAGATGAGG 961

Qy        901 CAGAGCCCAGCAAGCTGAATAGCATTGGCAATTACTACAAGCCGTGGTTCTTTAAGCATG 960
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Qy       1021 ACCACCGCCACACGCGCAGCATCTTCTGGGAGCTCCAGGACATCATCCCCTTTGGCAACA 1080
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Db       1082 ACCACCGCCACACGCGCAGCATCTTCTGGGAGCTCCAGGACATCATCCCCTTTGGCAACA 1141

Qy       1081 ACCCCATCTTCCGCTACCTCTTTGGCTGGATGGTGCCTCCCAAGATCTCCCTCCTGAAGC 1140
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Qy       1141 TGACCCAGGGTGAGACCCTGCGCAAGCTGTACGAGCAGCACCACGTGGTGCAGGACATGC 1200
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Qy       1201 TGGTGCCCATGAAGTGCCTGCAGCAGGCCCTGCACACCTTCCAAAACGACATCCACGTCT 1260
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Qy      1501 CCTTGTACCACAAGCTGCGAGAGAAGCTGGGTTGCCAGGACGCCTTCCCCGAGGTGTACG 1560
Db      1501 CCTTGTACCACAAGCTGCGAGAGAAGCTGGGTTGCCAGGACGCCTTCCCCGAGGTGTACG 1560
Qy      1561 ACAAGATCTGCAAGGCCGCCAGGCACTGAGCTGGAGCCCGCCTGGAGAGACAGACACGTG 1620
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Qy      1681 AAAGAGAAACCCCTCCAGAACTCCCACCCTGACAGCCCAACACCACCTTCCTCCTGGCTT 1740
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Qy      1741 CCAGGGGGCAGCCCAGTGGAATGGAAAGAATGTGGGATTTGGAGTCAGACAAGCCTGAGT 1800
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Qy      1801 CCAGTTCCCCGTTTAGAACTCATTAGCTGTGTGACTCTGGGTGAGTCCCTTAACCCCTCT 1860
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Qy      1861 GAGCCCGGGTCTCTTCATTAGTTGAAAGGGATAGTAATACCTACTTGCAGGTTGTTGTCA 1920
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Qy      1921 TCTGAGTTGAGCACTGGTCACATTGAAGGTGCTGGGTAAGTGGTAGCTCTTGTTGCTTCC 1980
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Qy      2041 AGCCATGGCTGGAATGATGAAGGGGATACGCTGGAGTTGCCCTGCCATCGCCTCCATCAG 2100
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Qy      2161 AGCCACGGAGTGGGGAGGCCCCAGATGCGCTGTGCCAAAGCCAGGTCCGAGGCCAAAGTT 2220
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Qy      2221 CTCCCTGCCATCCTTGGTGCCGTCCTGCCCCTTCCTCCTTCATGCCTGGGCCTGCAGGCC 2280
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Qy      2341 GGTTGAATCTTGTCCCAGCCTCAGCCTGGGACACCTAGGTGGAGAGAGTGGTCTCCGCTC 2400
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Qy      2401 TGAATTGGATCCAGGGGACCTGGGCTCATTCTTCTTGGCTCACCAACCCTGCAGGCCTCA 2460
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Qy      2581 CTCACCGTGGGGCTGGAAGAAAAGAGCATCGACTTGATTTCTCCAACCACTCATCCCTCT 2640
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Qy      2821 CCTCAGCTGAAAACCCGAATCTGAGAAGGAATTGCGTCATGTAAGGGAAGCTGGAATTAA 2880
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Qy      2881 GGGAGCTGAGCCAGTCATGGTTGTGGCGTGTGAGTCAGGAGACCTAGGTTTCAGCCCCTC 2940
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Qy      2941 TCTACTGTCAGCGAGCTGTGCAACGTGGGCAAGTCATTGTCCTCTGAGCTGCAGTTTCCT 3000
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Qy      3121 GTGGACACTTGGGGATTATCTGTCATTCAAGATCCTTCCTTCAACCCCAAGGCCAGCTCC 3180
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Qy      3181 CATCTCATTTCCAGAAAGGCTCATACCTGGCTTGCAGGGAAGCATCTGTCTTGTCATTCC 3240
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Qy      3241 AGGTGCCAGAATCCTCTCAGAGTCATTGAAGGGTGTTCACCCATCCCACCCAAGGCTTGG 3300
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Qy      3301 CACACTGCCAGTGTCTTAGCAGGGTCTTGTGAGGGCTGGGGGCATCCAGGCACTCAGAAG 3360
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Qy      3361 GCAAAGGAACCACCCTACCCATTTGGCCTCTGGAGGGGGCAGAAGAAAGAAAGAAACCTC 3420
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Qy      3421 ATCCTATATTTTACAAAGCATGTGAATTCTGGCATTAGCTCTCATAGGAGACCCATGTGC 3480
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Qy      3481 TTCCTTGCTCAGTGCAAAACTGATGATTCTACTTGCTGTAGATGAATGGTTAACACGAGC 3540
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RESULT 9
US-10-934-998-53
; Sequence 53, Application US/10934998
; Publication No. US20050153917A1
; GENERAL INFORMATION:
;  APPLICANT: AL-MAHMOOD, SALMAN
;  APPLICANT:  COLIN, SYLVIE
;  APPLICANT:  SCHNEIDER, CHRISTOPHE
;  TITLE OF INVENTION: GENES INVOLVED IN REGULATING ANGIOGENESIS, PHARMACEUTICAL
;  TITLE OF INVENTION:  PREPARATIONS CONTAINING SAME AND APPLICATIONS THEREOF
;  FILE REFERENCE: BMA-04-1206
;  CURRENT APPLICATION NUMBER: US/10/934,998
;  CURRENT FILING DATE:  2004-09-03
;  PRIOR APPLICATION NUMBER: PCT/FR03/00695
;  PRIOR FILING DATE: 2003-03-04
;  PRIOR APPLICATION NUMBER: FR02/02717
;  PRIOR FILING DATE: 2002-03-04
;  PRIOR APPLICATION NUMBER: FR02/04546
;  PRIOR FILING DATE: 2002-04-11
;  NUMBER OF SEQ ID NOS: 301
;  SOFTWARE: PatentIn version 3.2
; SEQ ID NO 53
;   LENGTH: 4187
;   TYPE: DNA
;   ORGANISM: Homo sapiens
;   FEATURE:
;   OTHER INFORMATION: GS-N53
;  PUBLICATION INFORMATION:
;   DATABASE ACCESSION NUMBER: GENBANK/D13643
;   DATABASE ENTRY DATE: 2001-10-06
;   RELEVANT RESIDUES: (1)..(4187)
US-10-934-998-53

  Query Match             100.0%;  Score 4187;  DB 22;  Length 4187;
  Best Local Similarity   100.0%;  Pred. No. 0;
  Matches 4187;  Conservative    0;  Mismatches    0;  Indels    0;  Gaps    0;

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Qy        301 CCTTCATGTGCACGGGGCGCCCTGGCTGGCTCACTGTCTCACTACGTGTCGGGAAGTACA 360
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Qy        661 CAGACCTGTTCTATGCCGTACCCTGGTCCTGTGGGACGCTGGGTTTCCTGGTGGCCGCTG 720
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Qy        901 CAGAGCCCAGCAAGCTGAATAGCATTGGCAATTACTACAAGCCGTGGTTCTTTAAGCATG 960
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Db       1141 TGACCCAGGGTGAGACCCTGCGCAAGCTGTACGAGCAGCACCACGTGGTGCAGGACATGC 1200

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Db       1201 TGGTGCCCATGAAGTGCCTGCAGCAGGCCCTGCACACCTTCCAAAACGACATCCACGTCT 1260

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Db       1261 ACCCCATCTGGCTGTGTCCGTTCATCCTGCCCAGCCAGCCAGGCCTAGTGCACCCCAAAG 1320

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Qy       1801 CCAGTTCCCCGTTTAGAACTCATTAGCTGTGTGACTCTGGGTGAGTCCCTTAACCCCTCT 1860
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Qy       1981 CGTTCAGCGTCACATCTGCAGTGGAGCCTGAAAAGGCTCCACATTAGGTCACCTGTGCAC 2040
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Db       1981 CGTTCAGCGTCACATCTGCAGTGGAGCCTGAAAAGGCTCCACATTAGGTCACCTGTGCAC 2040

Qy       2041 AGCCATGGCTGGAATGATGAAGGGGATACGCTGGAGTTGCCCTGCCATCGCCTCCATCAG 2100
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Qy       2101 CCAGACGAGGTCCTCACAGGAGAAGGACAGCTCTTCCCCACCCTGGGATCTCAGGAGGGC 2160
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Qy       2161 AGCCACGGAGTGGGGAGGCCCCAGATGCGCTGTGCCAAAGCCAGGTCCGAGGCCAAAGTT 2220
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Db       2161 AGCCACGGAGTGGGGAGGCCCCAGATGCGCTGTGCCAAAGCCAGGTCCGAGGCCAAAGTT 2220

Qy       2221 CTCCCTGCCATCCTTGGTGCCGTCCTGCCCCTTCCTCCTTCATGCCTGGGCCTGCAGGCC 2280
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Db       2221 CTCCCTGCCATCCTTGGTGCCGTCCTGCCCCTTCCTCCTTCATGCCTGGGCCTGCAGGCC 2280

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Qy       2341 GGTTGAATCTTGTCCCAGCCTCAGCCTGGGACACCTAGGTGGAGAGAGTGGTCTCCGCTC 2400
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Qy       2461 TCTTTCCCAAAACCCACTTTGTCTTGGTGGGAGTGGGTCCGCGCTGCTCTGCAGCAGGGG 2520
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Qy       2521 CTGGGGAGTGGACAGCATCAGGTGGGAAAGTGGAGTCCACCCTCATGTTTCTGTAGGATT 2580
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;  PRIOR FILING DATE: 2001-09-25
;  PRIOR APPLICATION NUMBER: US/09/962,832
;  PRIOR FILING DATE: 2001-09-25
;  PRIOR APPLICATION NUMBER: US/09/964,824
;  PRIOR FILING DATE: 2001-09-27
;  PRIOR APPLICATION NUMBER: US/09/967,768
;  PRIOR FILING DATE: 2001-09-28
;  PRIOR APPLICATION NUMBER: US/09/968,007
;  PRIOR FILING DATE: 2001-10-02
;  PRIOR APPLICATION NUMBER: US/09/969,347
;  PRIOR FILING DATE: 2001-10-02
;  PRIOR APPLICATION NUMBER: US/09/969,708
;  PRIOR FILING DATE: 2001-10-03
;  Remaining Prior Application data removed - See File Wrapper or PALM.
;  NUMBER OF SEQ ID NOS: 8447
;  SOFTWARE: PatentIn version 3.0
; SEQ ID NO 5139
;   LENGTH: 4187
;   TYPE: DNA
;   ORGANISM: Homo sapiens
US-10-843-641A-5139

  Query Match             100.0%;  Score 4187;  DB 21;  Length 4187;
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Db      3241 AGGTGCCAGAATCCTCTCAGAGTCATTGAAGGGTGTTCACCCATCCCACCCAAGGCTTGG 3300
Qy      3301 CACACTGCCAGTGTCTTAGCAGGGTCTTGTGAGGGCTGGGGGCATCCAGGCACTCAGAAG 3360
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Qy      3361 GCAAAGGAACCACCCTACCCATTTGGCCTCTGGAGGGGGCAGAAGAAAGAAAGAAACCTC 3420
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Qy      3421 ATCCTATATTTTACAAAGCATGTGAATTCTGGCATTAGCTCTCATAGGAGACCCATGTGC 3480
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Qy      3541 TAGTTAAACAGTGCCATTGTTTTGCCAGTGAAGCCTCCAACCCTAAGCCACTGGGACGGT 3600
Db      3541 TAGTTAAACAGTGCCATTGTTTTGCCAGTGAAGCCTCCAACCCTAAGCCACTGGGACGGT 3600
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Qy      3901 CCTGAGTTTCTTGGCTGCATGAAGGTTTTCTTTAGAATCAGAATTGTAGCCAGTTTCTTT 3960
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Qy      3961 GGCCAGAAGGATGAATACTTGGATATTACTGAAAGGGAGGGGTGGAGATGGGTGTGGCAG 4020
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Db      4081 ATCTTCCCTGTCAGGCTCTTACAGCCACAGGCACTGTGTCTACTGTCTGGAAGACATGTC 4140
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Db      4141 CCCGTGGCTGTGGGGCCGCTGCTTCTGTTTAAATAAAAGTGGCCTGG 4187
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RESULT 8
US-10-843-641A-5139
; Sequence 5139, Application US/10843641A
; Publication No. US20050064454A1
; GENERAL INFORMATION:
;  APPLICANT: Avalon Pharmaceuticals, Inc.
;  TITLE OF INVENTION: Cancer Gene Determination and Therapeutic Screening Using
;  TITLE OF INVENTION:  Signature Gene Sets
;  FILE REFERENCE: 689290-189
;  CURRENT APPLICATION NUMBER: US/10/843,641A
;  CURRENT FILING DATE:  2004-05-12
;  PRIOR APPLICATION NUMBER: US/09/873,367
;  PRIOR FILING DATE: 2001-06-05
;  PRIOR APPLICATION NUMBER: US/09/954,531
;  PRIOR FILING DATE: 2001-09-18
;  PRIOR APPLICATION NUMBER: US/09/954,456
;  PRIOR FILING DATE: 2001-09-25
;  PRIOR APPLICATION NUMBER: US/09/962,436
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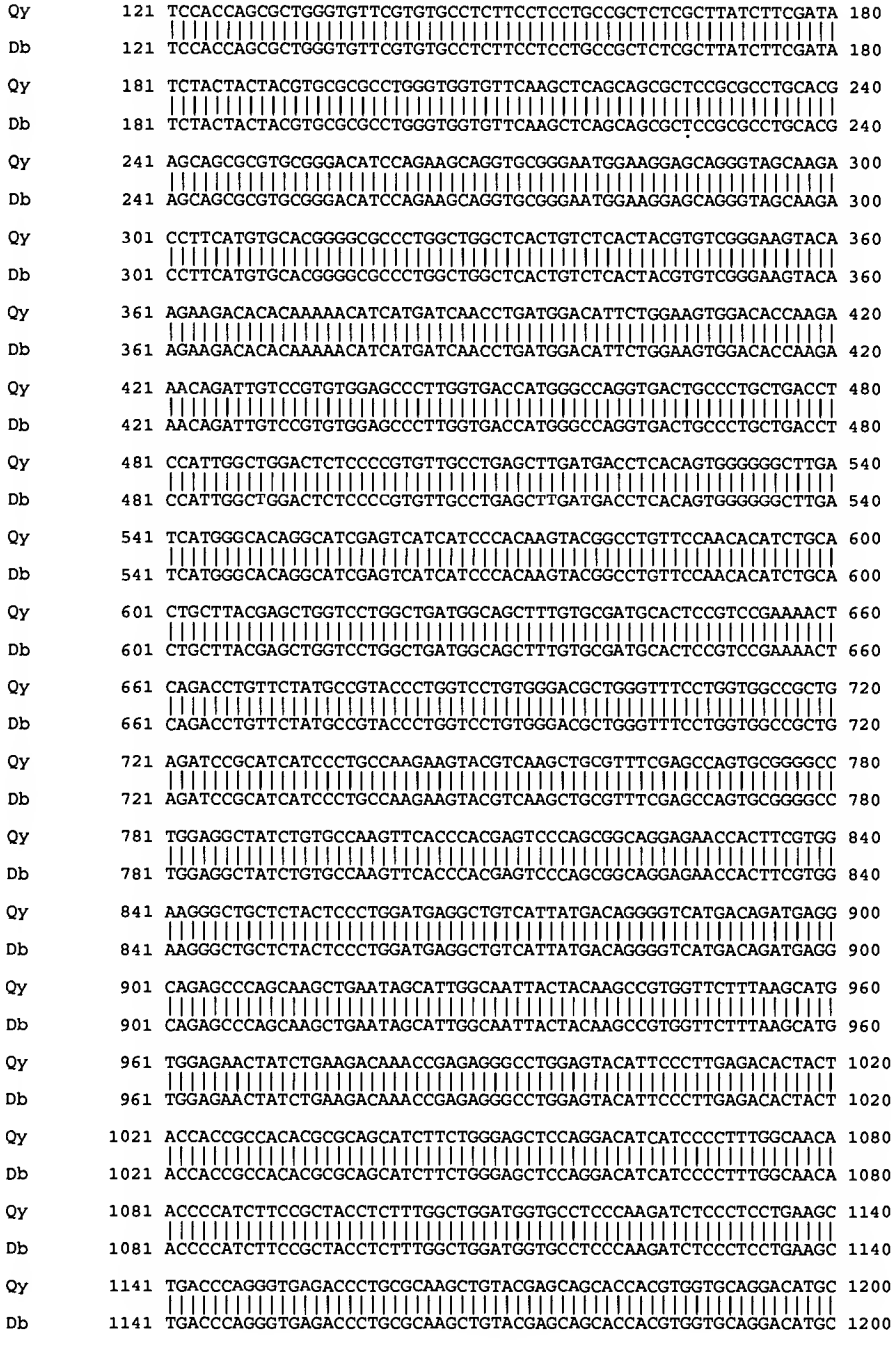
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Db 361 AGAAGACACACAAAAACATCATGATCAACCTGATGGACATTCTGGAAGTGGACACCAAGA 420
Qy 421 AACAGATTGTCCGTGTGGAGCCCTTGGTGACCATGGGCCAGGTGACTGCCCTGCTGACCT 480
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Db 661 CAGACCTGTTCTATGCCGTACCCTGGTCCTGTGGGACGCTGGGTTTCCTGGTGGCCGCTG 720
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Db 721 AGATCCGCATCATCCCTGCCAAGAAGTACGTCAAGCTGCGTTTCGAGCCAGTGCGGGGCC 780
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Qy 841 AAGGGCTGCTCTACTCCCTGGATGAGGCTGTCATTATGACAGGGGTCATGACAGATGAGG 900
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Qy 901 CAGAGCCCAGCAAGCTGAATAGCATTGGCAATTACTACAAGCCGTGGTTCTTTAAGCATG 960
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Qy 1201 TGGTGCCCATGAAGTGCCTGCAGCAGGCCCTGCACACCTTCCAAAACGACATCCACGTCT 1260
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Qy 1921 TCTGAGTTGAGCACTGGTCACATTGAAGGTGCTGGGTAAGTGGTAGCTCTTGTTGCTTCC 1980
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Qy 2041 AGCCATGGCTGGAATGATGAAGGGGATACGCTGGAGTTGCCCTGCCATCGCCTCCATCAG 2100
Db 2041 AGCCATGGCTGGAATGATGAAGGGGATACGCTGGAGTTGCCCTGCCATCGCCTCCATCAG 2100
Qy 2101 CCAGACGAGGTCCTCACAGGAGAAGGACAGCTCTTCCCCACCCTGGGATCTCAGGAGGGC 2160
Db 2101 CCAGACGAGGTCCTCACAGGAGAAGGACAGCTCTTCCCCACCCTGGGATCTCAGGAGGGC 2160
Qy 2161 AGCCACGGAGTGGGGAGGCCCCAGATGCGCTGTGCCAAAGCCAGGTCCGAGGCCAAAGTT 2220
Db 2161 AGCCACGGAGTGGGGAGGCCCCAGATGCGCTGTGCCAAAGCCAGGTCCGAGGCCAAAGTT 2220
Qy 2221 CTCCCTGCCATCCTTGGTGCCGTCCTGCCCCTTCCTCCTTCATGCCTGGGCCTGCAGGCC 2280
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Qy 2281 CACCCCAGCCACCACTGAGTCCACTCGGAGTGCCCTGTGTTCCTGGAGAAGGCATTCCAG 2340

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Db       2641 TTTTCTTTCTTCCACCACTCCCCACCCCAGCTGTAGTTAATTTCAGTGCCTTACAAATCC 2700

Qy       2701 TAAGCTCAGAGAAAGTTCCATTTCCGTTCCAGAGGGAAGGGAACCTCCCTAGGTCCTTCC 2760
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Db       2701 TAAGCTCAGAGAAAGTTCCATTTCCGTTCCAGAGGGAAGGGAACCTCCCTAGGTCCTTCC 2760

Qy       2761 CTGGCTTGTTATAACGCAAAGCTTGGTTGTTTATGCAACTCTATCTTAAGAACTGCCCAG 2820
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Db       2761 CTGGCTTGTTATAACGCAAAGCTTGGTTGTTTATGCAACTCTATCTTAAGAACTGCCCAG 2820

Qy       2821 CCTCAGCTGAAAACCCGAATCTGAGAAGGAATTGCGTCATGTAAGGGAAGCTGGAATTAA 2880
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Qy       3181 CATCTCATTTCCAGAAAGGCTCATACCTGGCTTGCAGGGAAGCATCTGTCTTGTCATTCC 3240
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Qy       3841 CTTGATGCCAGTGGGCAGTGCATGTGGAAAGTATCAGAGTAAGCCTCTCCCCCTCCAGAGC 3900
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RESULT 7
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; Sequence 355, Application US/10342887
; Publication No. US20040058340A1
; GENERAL INFORMATION:
; APPLICANT: Dai, Hongyue
; APPLICANT: He, Yudong
; APPLICANT: Linsley, Peter S.
; APPLICANT: Mao, Mao
; APPLICANT: Roberts, Christopher J.
; APPLICANT: Van 't Veer, Laura Johanna
; APPLICANT: Van de Vijver, Marc J.
; APPLICANT: Bernards, Rene
; TITLE OF INVENTION: Diagnosis and Prognosis of Breast Cancer Patients
; FILE REFERENCE: 9301-188-999
; CURRENT APPLICATION NUMBER: US/10/342,887
; CURRENT FILING DATE: 2003-01-15
; PRIOR APPLICATION NUMBER: 60/298,918
; PRIOR FILING DATE: 2001-06-18
; PRIOR APPLICATION NUMBER: 60/380,710
; PRIOR FILING DATE: 2002-05-14
; PRIOR APPLICATION NUMBER: 10/172,118
; PRIOR FILING DATE: 2002-06-14
; NUMBER OF SEQ ID NOS: 2699
; SEQ ID NO 355
; LENGTH: 4187
; TYPE: DNA
; ORGANISM: Homo sapiens
US-10-342-887-355

Query Match 100.0%; Score 4187; DB 18; Length 4187;
Best Local Similarity 100.0%; Pred. No. 0;
Matches 4187; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

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Qy        661 CAGACCTGTTCTATGCCGTACCCTGGTCCTGTGGGACGCTGGGTTTCCTGGTGGCCGCTG 720
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Qy        721 AGATCCGCATCATCCCTGCCAAGAAGTACGTCAAGCTGCGTTTCGAGCCAGTGCGGGGCC 780
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Qy        841 AAGGGCTGCTCTACTCCCTGGATGAGGCTGTCATTATGACAGGGGTCATGACAGATGAGG 900
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Qy        961 TGGAGAACTATCTGAAGACAAACCGAGAGGGCCTGGAGTACATTCCCTTGAGACACTACT 1020
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Qy       1081 ACCCCATCTTCCGCTACCTCTTTGGCTGGATGGTGCCTCCCAAGATCTCCCTCCTGAAGC 1140
Db       1081 ACCCCATCTTCCGCTACCTCTTTGGCTGGATGGTGCCTCCCAAGATCTCCCTCCTGAAGC 1140
Qy       1141 TGACCCAGGGTGAGACCCTGCGCAAGCTGTACGAGCAGCACCACGTGGTGCAGGACATGC 1200
Db       1141 TGACCCAGGGTGAGACCCTGCGCAAGCTGTACGAGCAGCACCACGTGGTGCAGGACATGC 1200
Qy       1201 TGGTGCCCATGAAGTGCCTGCAGCAGGCCCTGCACACCTTCCAAAACGACATCCACGTCT 1260
Db       1201 TGGTGCCCATGAAGTGCCTGCAGCAGGCCCTGCACACCTTCCAAAACGACATCCACGTCT 1260
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Qy       1321 GAAATGAGGCAGAGCTCTACATCGACATTGGAGCATATGGGGAGCCGCGTGTGAAACACT 1380
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Qy       1381 TTGAAGCCAGGTCCTGCATGAGGCAGCTGGAGAAGTTTGTCCGCAGCGTGCATGGCTTCC 1440
Db       1381 TTGAAGCCAGGTCCTGCATGAGGCAGCTGGAGAAGTTTGTCCGCAGCGTGCATGGCTTCC 1440
Qy       1441 AGATGCTGTATGCCGACTGCTACATGAACCGGGAGGAGTTCTGGGAGATGTTTGATGGCT 1500
Db       1441 AGATGCTGTATGCCGACTGCTACATGAACCGGGAGGAGTTCTGGGAGATGTTTGATGGCT 1500
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Db       1621 TGAGTGGTCAGGCATCTTCCCTTCACTCAAGCTTGGCTGCTTTCCTAGATCCACACTTTC 1680
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Db       1681 AAAGAGAAACCCCTCCAGAACTCCCACCCTGACAGCCCAACACCACCTTCCTCCTGGCTT 1740
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Db       1741 CCAGGGGGCAGCCCAGTGGAATGGAAAGAATGTGGGATTTGGAGTCAGACAAGCCTGAGT 1800
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Db       1801 CCAGTTCCCCGTTTAGAACTCATTAGCTGTGTGACTCTGGGTGAGTCCCTTAACCCCTCT 1860
Qy       1861 GAGCCCGGGTCTCTTCATTAGTTGAAAGGGATAGTAATACCTACTTGCAGGTTGTTGTCA 1920
Db       1861 GAGCCCGGGTCTCTTCATTAGTTGAAAGGGATAGTAATACCTACTTGCAGGTTGTTGTCA 1920
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Qy       2041 AGCCATGGCTGGAATGATGAAGGGGATACGCTGGAGTTGCCCTGCCATCGCCTCCATCAG 2100
Db       2041 AGCCATGGCTGGAATGATGAAGGGGATACGCTGGAGTTGCCCTGCCATCGCCTCCATCAG 2100
Qy       2101 CCAGACGAGGTCCTCACAGGAGAAGGACAGCTCTTCCCCACCCTGGGATCTCAGGAGGGC 2160
Db       2101 CCAGACGAGGTCCTCACAGGAGAAGGACAGCTCTTCCCCACCCTGGGATCTCAGGAGGGC 2160
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Db       2161 AGCCACGGAGTGGGGAGGCCCCAGATGCGCTGTGCCAAAGCCAGGTCCGAGGCCAAAGTT 2220
Qy       2221 CTCCCTGCCATCCTTGGTGCCGTCCTGCCCCTTCCTCCTTCATGCCTGGGCCTGCAGGCC 2280
Db       2221 CTCCCTGCCATCCTTGGTGCCGTCCTGCCCCTTCCTCCTTCATGCCTGGGCCTGCAGGCC 2280
Qy       2281 CACCCCAGCCACCACTGAGTCCACTCGGAGTGCCCTGTGTTCCTGGAGAAGGCATTCCAG 2340
Db       2281 CACCCCAGCCACCACTGAGTCCACTCGGAGTGCCCTGTGTTCCTGGAGAAGGCATTCCAG 2340
Qy       2341 GGTTGAATCTTGTCCCAGCCTCAGCCTGGGACACCTAGGTGGAGAGAGTGGTCTCCGCTC 2400
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Qy       2401 TGAATTGGATCCAGGGGACCTGGGCTCATTCTTCTTGGCTCACCAACCCTGCAGGCCTCA 2460
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Qy       2461 TCTTTCCCAAAACCCACTTTGTCTTGGTGGGAGTGGGTCCGCGCTGCTCTGCAGCAGGGG 2520
Db       2461 TCTTTCCCAAAACCCACTTTGTCTTGGTGGGAGTGGGTCCGCGCTGCTCTGCAGCAGGGG 2520
Qy       2521 CTGGGGAGTGGACAGCATCAGGTGGGAAAGTGGAGTCCACCCTCATGTTTCTGTAGGATT 2580
Db       2521 CTGGGGAGTGGACAGCATCAGGTGGGAAAGTGGAGTCCACCCTCATGTTTCTGTAGGATT 2580
Qy       2581 CTCACCGTGGGGCTGGAAGAAAAGAGCATCGACTTGATTTCTCCAACCACTCATCCCTCT 2640
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Qy       3001 CATCTGTCACATCGCTACAGACAAGACCTCCCTGGAACCCTTCTGATTGTCTTAGACACT 3060
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Qy       3121 GTGGACACTTGGGGATTATCTGTCATTCAAGATCCTTCCTTCAACCCCAAGGCCAGCTCC 3180
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Db       3121 GTGGACACTTGGGGATTATCTGTCATTCAAGATCCTTCCTTCAACCCCAAGGCCAGCTCC 3180

Qy       3181 CATCTCATTTCCAGAAAGGCTCATACCTGGCTTGCAGGGAAGCATCTGTCTTGTCATTCC 3240
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Db       3181 CATCTCATTTCCAGAAAGGCTCATACCTGGCTTGCAGGGAAGCATCTGTCTTGTCATTCC 3240

Qy       3241 AGGTGCCAGAATCCTCTCAGAGTCATTGAAGGGTGTTCACCCATCCCACCCAAGGCTTGG 3300
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Db       3241 AGGTGCCAGAATCCTCTCAGAGTCATTGAAGGGTGTTCACCCATCCCACCCAAGGCTTGG 3300

Qy       3301 CACACTGCCAGTGTCTTAGCAGGGTCTTGTGAGGGCTGGGGGCATCCAGGCACTCAGAAG 3360
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Qy       3421 ATCCTATATTTTACAAAGCATGTGAATTCTGGCATTAGCTCTCATAGGAGACCCATGTGC 3480
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Db       3421 ATCCTATATTTTACAAAGCATGTGAATTCTGGCATTAGCTCTCATAGGAGACCCATGTGC 3480

Qy       3481 TTCCTTGCTCAGTGCAAAACTGATGATTCTACTTGCTGTAGATGAATGGTTAACACGAGC 3540
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Db       3481 TTCCTTGCTCAGTGCAAAACTGATGATTCTACTTGCTGTAGATGAATGGTTAACACGAGC 3540

Qy       3541 TAGTTAAACAGTGCCATTGTTTTGCCAGTGAAGCCTCCAACCCTAAGCCACTGGGACGGT 3600
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Qy       3601 GGCCAGAGATGCCAGCAGCCTCTGTCGCCCTTAGTCATATAACCAAAATCCAGACCTTAT 3660
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Qy       3661 CCACAACCCGGGGCTTGGAAAGGAAGGTATTTTGGAATCACACCCTCCGGTTATGTTGCT 3720
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       3661 CCACAACCCGGGGCTTGGAAAGGAAGGTATTTTGGAATCACACCCTCCGGTTATGTTGCT 3720

Qy       3721 CCAGTAAAATCTTGCCTGGAAAGAGGCAGTCTTCTTAGCATGGTGAGCTGAGTTCATGGC 3780
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       3721 CCAGTAAAATCTTGCCTGGAAAGAGGCAGTCTTCTTAGCATGGTGAGCTGAGTTCATGGC 3780

Qy       3781 TTTTTTTTGTAGCCAGTCCTGTCCCTGGCCATCCATGTGATGGTTTTGGATGGAGTTAAA 3840
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Db       3781 TTTTTTTTGTAGCCAGTCCTGTCCCTGGCCATCCATGTGATGGTTTTGGATGGAGTTAAA 3840

Qy       3841 CTTGATGCCAGTGGGCAGTGCATGTGGAAAGTATCAGAGTAAGCCTCTCCCCTCCAGAGC 3900
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       3841 CTTGATGCCAGTGGGCAGTGCATGTGGAAAGTATCAGAGTAAGCCTCTCCCCTCCAGAGC 3900

Qy       3901 CCTGAGTTTCTTGGCTGCATGAAGGTTTTCTTTAGAATCAGAATTGTAGCCAGTTTCTTT 3960
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Qy       4021 TGTATGGTGTGTGATTTTTATTTTCTTCTTTGGTCATGGGGGCCAAGGAGAAAGGCATGA 4080
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Qy       4141 CCCGTGGCTGTGGGGCCGCTGCTTCTGTTTAAATAAAAGTGGCCTGG 4187
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RESULT 6
US-10-240-425-1203
; Sequence 1203, Application US/10240425
; Publication No. US20040033502A1
; GENERAL INFORMATION:
; APPLICANT: Williams, Amanda
; APPLICANT: Boland, Joseph F.
; APPLICANT: Lord, Reginald V.
; APPLICANT: Alvarez, Chris
; APPLICANT: Wetzel, Jon C.
; APPLICANT: Scherf, Uwe
; APPLICANT: Vockley, Joseph G.
; TITLE OF INVENTION: Gene Expression Profiles in Esophageal Tissue
; FILE REFERENCE: 44921-5026
; CURRENT APPLICATION NUMBER: US/10/240,425
; CURRENT FILING DATE: 2002-09-30
; PRIOR APPLICATION NUMBER: PCT/US01/09847
; PRIOR FILING DATE: 2001-03-28
; PRIOR APPLICATION NUMBER: US 60/193,446
; PRIOR FILING DATE: 2000-03-31
; NUMBER OF SEQ ID NOS: 1588
; SOFTWARE: PatentIn Ver. 2.1
; SEQ ID NO 1203
; LENGTH: 4187
; TYPE: DNA
; ORGANISM: Homo sapiens
; FEATURE:
; OTHER INFORMATION: Genbank Accession No. US20040033502A1 D13643
US-10-240-425-1203

Query Match 100.0%; Score 4187; DB 18; Length 4187;
Best Local Similarity 100.0%; Pred. No. 0;
Matches 4187; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

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Qy        121 TCCACCAGCGCTGGGTGTTCGTGTGCCTCTTCCTCCTGCCGCTCTCGCTTATCTTCGATA 180
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Qy        181 TCTACTACTACGTGCGCGCCTGGGTGGTGTTCAAGCTCAGCAGCGCTCCGCGCCTGCACG 240
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Qy        301 CCTTCATGTGCACGGGGCGCCCTGGCTGGCTCACTGTCTCACTACGTGTCGGGAAGTACA 360
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Qy        361 AGAAGACACACAAAAACATCATGATCAACCTGATGGACATTCTGGAAGTGGACACCAAGA 420
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Qy        421 AACAGATTGTCCGTGTGGAGCCCTTGGTGACCATGGGCCAGGTGACTGCCCTGCTGACCT 480
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Qy        901 CAGAGCCCAGCAAGCTGAATAGCATTGGCAATTACTACAAGCCGTGGTTCTTTAAGCATG 960
Db        901 CAGAGCCCAGCAAGCTGAATAGCATTGGCAATTACTACAAGCCGTGGTTCTTTAAGCATG 960
Qy        961 TGGAGAACTATCTGAAGACAAACCGAGAGGGCCTGGAGTACATTCCCTTGAGACACTACT 1020
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Qy       2401 TGAATTGGATCCAGGGGACCTGGGCTCATTCTTCTTGGCTCACCAACCCTGCAGGCCTCA 2460
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Qy       2641 TTTTCTTTCTTCCACCACTCCCCACCCCAGCTGTAGTTAATTTCAGTGCCTTACAAATCC 2700
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Qy       3481 TTCCTTGCTCAGTGCAAAACTGATGATTCTACTTGCTGTAGATGAATGGTTAACACGAGC 3540
                ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       3481 TTCCTTGCTCAGTGCAAAACTGATGATTCTACTTGCTGTAGATGAATGGTTAACACGAGC 3540

Qy       3541 TAGTTAAACAGTGCCATTGTTTTGCCAGTGAAGCCTCCAACCCTAAGCCACTGGGACGGT 3600
                ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       3541 TAGTTAAACAGTGCCATTGTTTTGCCAGTGAAGCCTCCAACCCTAAGCCACTGGGACGGT 3600

Qy       3601 GGCCAGAGATGCCAGCAGCCTCTGTCGCCCTTAGTCATATAACCAAAATCCAGACCTTAT 3660
                ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       3601 GGCCAGAGATGCCAGCAGCCTCTGTCGCCCTTAGTCATATAACCAAAATCCAGACCTTAT 3660

Qy       3661 CCACAACCCGGGGCTTGGAAAGGAAGGTATTTTGGAATCACACCCTCCGGTTATGTTGCT 3720
                ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       3661 CCACAACCCGGGGCTTGGAAAGGAAGGTATTTTGGAATCACACCCTCCGGTTATGTTGCT 3720

Qy       3721 CCAGTAAAATCTTGCCTGGAAAGAGGCAGTCTTCTTAGCATGGTGAGCTGAGTTCATGGC 3780
                ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       3721 CCAGTAAAATCTTGCCTGGAAAGAGGCAGTCTTCTTAGCATGGTGAGCTGAGTTCATGGC 3780

Qy       3781 TTTTTTTTGTAGCCAGTCCTGTCCCTGGCCATCCATGTGATGGTTTTGGATGGAGTTAAA 3840
                ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       3781 TTTTTTTTGTAGCCAGTCCTGTCCCTGGCCATCCATGTGATGGTTTTGGATGGAGTTAAA 3840

Qy       3841 CTTGATGCCAGTGGGCAGTGCATGTGGAAAGTATCAGAGTAAGCCTCTCCCCTCCAGAGC 3900
                ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       3841 CTTGATGCCAGTGGGCAGTGCATGTGGAAAGTATCAGAGTAAGCCTCTCCCCTCCAGAGC 3900

Qy       3901 CCTGAGTTTCTTGGCTGCATGAAGGTTTTCTTTAGAATCAGAATTGTAGCCAGTTTCTTT 3960
                ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       3901 CCTGAGTTTCTTGGCTGCATGAAGGTTTTCTTTAGAATCAGAATTGTAGCCAGTTTCTTT 3960

Qy       3961 GGCCAGAAGGATGAATACTTGGATATTACTGAAAGGGAGGGGTGGAGATGGGTGTGGCAG 4020
                ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       3961 GGCCAGAAGGATGAATACTTGGATATTACTGAAAGGGAGGGGTGGAGATGGGTGTGGCAG 4020

Qy       4021 TGTATGGTGTGTGATTTTTATTTTCTTCTTTGGTCATGGGGGCCAAGGAGAAAGGCATGA 4080
                ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       4021 TGTATGGTGTGTGATTTTTATTTTCTTCTTTGGTCATGGGGGCCAAGGAGAAAGGCATGA 4080

Qy       4081 ATCTTCCCTGTCAGGCTCTTACAGCCACAGGCACTGTGTCTACTGTCTGGAAGACATGTC 4140
                ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       4081 ATCTTCCCTGTCAGGCTCTTACAGCCACAGGCACTGTGTCTACTGTCTGGAAGACATGTC 4140

Qy       4141 CCCGTGGCTGTGGGGCCGCTGCTTCTGTTTAAATAAAAGTGGCCTGG 4187
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Db       4141 CCCGTGGCTGTGGGGCCGCTGCTTCTGTTTAAATAAAAGTGGCCTGG 4187


RESULT 5
US-10-172-118-355
; Sequence 355, Application US/10172118
; Publication No. US20030224374A1
; GENERAL INFORMATION:
;  APPLICANT: Dai, Hongyue
;  APPLICANT:  He, Yudong
;  APPLICANT:  Linsley, Peter
;  APPLICANT:  Mao, Mao
;  APPLICANT:  Roberts, Chris
;  APPLICANT:  Van 't Veer, Laura
;  APPLICANT:  Van de Vijver, Marc
;  APPLICANT:  Bernards, Rene
;  TITLE OF INVENTION: Diagnosis and Prognosis of Breast Cancer Patients
;  FILE REFERENCE: 9301-175-999
;  CURRENT APPLICATION NUMBER: US/10/172,118
;  CURRENT FILING DATE:  2002-06-14
;  PRIOR APPLICATION NUMBER: 60/380,770
;  PRIOR FILING DATE: 2002-05-14
;  NUMBER OF SEQ ID NOS: 2699
; SEQ ID NO 355
;   LENGTH: 4187
;   TYPE: DNA
;   ORGANISM: Homo sapiens
;  PUBLICATION INFORMATION:
;   DATABASE ACCESSION NUMBER: D13643
;   DATABASE ENTRY DATE: 2001-06-18
US-10-172-118-355

  Query Match             100.0%;  Score 4187;  DB 17;  Length 4187;
  Best Local Similarity   100.0%;  Pred. No. 0;
  Matches 4187;  Conservative    0;  Mismatches    0;  Indels    0;  Gaps    0;

Qy          1 GGCGCGAACCCGCAGCGCTTACCGCGCGGCGCCGCACCATGGAGCCCGCCGTGTCGCTGG 60
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db          1 GGCGCGAACCCGCAGCGCTTACCGCGCGGCGCCGCACCATGGAGCCCGCCGTGTCGCTGG 60

Qy         61 CCGTGTGCGCGCTGCTCTTCCTGCTGTGGGTGCGCCTGAAGGGGCTGGAGTTCGTGCTCA 120
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db         61 CCGTGTGCGCGCTGCTCTTCCTGCTGTGGGTGCGCCTGAAGGGGCTGGAGTTCGTGCTCA 120

Qy        121 TCCACCAGCGCTGGGTGTTCGTGTGCCTCTTCCTCCTGCCGCTCTCGCTTATCTTCGATA 180
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        121 TCCACCAGCGCTGGGTGTTCGTGTGCCTCTTCCTCCTGCCGCTCTCGCTTATCTTCGATA 180

Qy        181 TCTACTACTACGTGCGCGCCTGGGTGGTGTTCAAGCTCAGCAGCGCTCCGCGCCTGCACG 240
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        181 TCTACTACTACGTGCGCGCCTGGGTGGTGTTCAAGCTCAGCAGCGCTCCGCGCCTGCACG 240

Qy        241 AGCAGCGCGTGCGGGACATCCAGAAGCAGGTGCGGGAATGGAAGGAGCAGGGTAGCAAGA 300
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        241 AGCAGCGCGTGCGGGACATCCAGAAGCAGGTGCGGGAATGGAAGGAGCAGGGTAGCAAGA 300

Qy        301 CCTTCATGTGCACGGGGCGCCCTGGCTGGCTCACTGTCTCACTACGTGTCGGGAAGTACA 360
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        301 CCTTCATGTGCACGGGGCGCCCTGGCTGGCTCACTGTCTCACTACGTGTCGGGAAGTACA 360

Qy        361 AGAAGACACACAAAAACATCATGATCAACCTGATGGACATTCTGGAAGTGGACACCAAGA 420
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        361 AGAAGACACACAAAAACATCATGATCAACCTGATGGACATTCTGGAAGTGGACACCAAGA 420

Qy        421 AACAGATTGTCCGTGTGGAGCCCTTGGTGACCATGGGCCAGGTGACTGCCCTGCTGACCT 480
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        421 AACAGATTGTCCGTGTGGAGCCCTTGGTGACCATGGGCCAGGTGACTGCCCTGCTGACCT 480

Qy        481 CCATTGGCTGGACTCTCCCCGTGTTGCCTGAGCTTGATGACCTCACAGTGGGGGGCTTGA 540
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        481 CCATTGGCTGGACTCTCCCCGTGTTGCCTGAGCTTGATGACCTCACAGTGGGGGGCTTGA 540

Qy        541 TCATGGGCACAGGCATCGAGTCATCATCCCACAAGTACGGCCTGTTCCAACACATCTGCA 600
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        541 TCATGGGCACAGGCATCGAGTCATCATCCCACAAGTACGGCCTGTTCCAACACATCTGCA 600

Qy        601 CTGCTTACGAGCTGGTCCTGGCTGATGGCAGCTTTGTGCGATGCACTCCGTCCGAAAACT 660
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        601 CTGCTTACGAGCTGGTCCTGGCTGATGGCAGCTTTGTGCGATGCACTCCGTCCGAAAACT 660

Qy        661 CAGACCTGTTCTATGCCGTACCCTGGTCCTGTGGGACGCTGGGTTTCCTGGTGGCCGCTG 720
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        661 CAGACCTGTTCTATGCCGTACCCTGGTCCTGTGGGACGCTGGGTTTCCTGGTGGCCGCTG 720

Qy        721 AGATCCGCATCATCCCTGCCAAGAAGTACGTCAAGCTGCGTTTCGAGCCAGTGCGGGGCC 780
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        721 AGATCCGCATCATCCCTGCCAAGAAGTACGTCAAGCTGCGTTTCGAGCCAGTGCGGGGCC 780

Qy        781 TGGAGGCTATCTGTGCCAAGTTCACCCACGAGTCCCAGCGGCAGGAGAACCACTTCGTGG 840
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Qy      1141 TGACCCAGGGTGAGACCCTGCGCAAGCTGTACGAGCAGCACCACGTGGTGCAGGACATGC 1200
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db      1141 TGACCCAGGGTGAGACCCTGCGCAAGCTGTACGAGCAGCACCACGTGGTGCAGGACATGC 1200

Qy      1201 TGGTGCCCATGAAGTGCCTGCAGCAGGCCCTGCACACCTTCCAAAACGACATCCACGTCT 1260
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db      1201 TGGTGCCCATGAAGTGCCTGCAGCAGGCCCTGCACACCTTCCAAAACGACATCCACGTCT 1260

Qy      1261 ACCCCATCTGGCTGTGTCCGTTCATCCTGCCCAGCCAGCCAGGCCTAGTGCACCCCAAAG 1320
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db      1261 ACCCCATCTGGCTGTGTCCGTTCATCCTGCCCAGCCAGCCAGGCCTAGTGCACCCCAAAG 1320

Qy      1321 GAAATGAGGCAGAGCTCTACATCGACATTGGAGCATATGGGGAGCCGCGTGTGAAACACT 1380
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db      1321 GAAATGAGGCAGAGCTCTACATCGACATTGGAGCATATGGGGAGCCGCGTGTGAAACACT 1380

Qy      1381 TTGAAGCCAGGTCCTGCATGAGGCAGCTGGAGAAGTTTGTCCGCAGCGTGCATGGCTTCC 1440
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db      1381 TTGAAGCCAGGTCCTGCATGAGGCAGCTGGAGAAGTTTGTCCGCAGCGTGCATGGCTTCC 1440

Qy      1441 AGATGCTGTATGCCGACTGCTACATGAACCGGGAGGAGTTCTGGGAGATGTTTGATGGCT 1500
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db      1441 AGATGCTGTATGCCGACTGCTACATGAACCGGGAGGAGTTCTGGGAGATGTTTGATGGCT 1500

Qy      1501 CCTTGTACCACAAGCTGCGAGAGAAGCTGGGTTGCCAGGACGCCTTCCCCGAGGTGTACG 1560
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db      1501 CCTTGTACCACAAGCTGCGAGAGAAGCTGGGTTGCCAGGACGCCTTCCCCGAGGTGTACG 1560

Qy      1561 ACAAGATCTGCAAGGCCGCCAGGCACTGAGCTGGAGCCCGCCTGGAGAGACAGACACGTG 1620
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db      1561 ACAAGATCTGCAAGGCCGCCAGGCACTGAGCTGGAGCCCGCCTGGAGAGACAGACACGTG 1620

Qy      1621 TGAGTGGTCAGGCATCTTCCCTTCACTCAAGCTTGGCTGCTTTCCTAGATCCACACTTTC 1680
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db      1621 TGAGTGGTCAGGCATCTTCCCTTCACTCAAGCTTGGCTGCTTTCCTAGATCCACACTTTC 1680

Qy      1681 AAAGAGAAACCCCTCCAGAACTCCCACCCTGACAGCCCAACACCACCTTCCTCCTGGCTT 1740
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db      1681 AAAGAGAAACCCCTCCAGAACTCCCACCCTGACAGCCCAACACCACCTTCCTCCTGGCTT 1740

Qy      1741 CCAGGGGGCAGCCCAGTGGAATGGAAAGAATGTGGGATTTGGAGTCAGACAAGCCTGAGT 1800
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db      1741 CCAGGGGGCAGCCCAGTGGAATGGAAAGAATGTGGGATTTGGAGTCAGACAAGCCTGAGT 1800

Qy      1801 CCAGTTCCCCGTTTAGAACTCATTAGCTGTGTGACTCTGGGTGAGTCCCTTAACCCCTCT 1860
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db      1801 CCAGTTCCCCGTTTAGAACTCATTAGCTGTGTGACTCTGGGTGAGTCCCTTAACCCCTCT 1860

Qy      1861 GAGCCCGGGTCTCTTCATTAGTTGAAAGGGATAGTAATACCTACTTGCAGGTTGTTGTCA 1920
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db      1861 GAGCCCGGGTCTCTTCATTAGTTGAAAGGGATAGTAATACCTACTTGCAGGTTGTTGTCA 1920

Qy      1921 TCTGAGTTGAGCACTGGTCACATTGAAGGTGCTGGGTAAGTGGTAGCTCTTGTTGCTTCC 1980
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db      1921 TCTGAGTTGAGCACTGGTCACATTGAAGGTGCTGGGTAAGTGGTAGCTCTTGTTGCTTCC 1980

Qy      1981 CGTTCAGCGTCACATCTGCAGTGGAGCCTGAAAAGGCTCCACATTAGGTCACCTGTGCAC 2040
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db      1981 CGTTCAGCGTCACATCTGCAGTGGAGCCTGAAAAGGCTCCACATTAGGTCACCTGTGCAC 2040

Qy      2041 AGCCATGGCTGGAATGATGAAGGGGATACGCTGGAGTTGCCCTGCCATCGCCTCCATCAG 2100
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db      2041 AGCCATGGCTGGAATGATGAAGGGGATACGCTGGAGTTGCCCTGCCATCGCCTCCATCAG 2100

Qy      2101 CCAGACGAGGTCCTCACAGGAGAAGGACAGCTCTTCCCCACCCTGGGATCTCAGGAGGGC 2160
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db      2101 CCAGACGAGGTCCTCACAGGAGAAGGACAGCTCTTCCCCACCCTGGGATCTCAGGAGGGC 2160

Qy      2161 AGCCACGGAGTGGGGAGGCCCCAGATGCGCTGTGCCAAAGCCAGGTCCGAGGCCAAAGTT 2220
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db      2161 AGCCACGGAGTGGGGAGGCCCCAGATGCGCTGTGCCAAAGCCAGGTCCGAGGCCAAAGTT 2220

Qy      2221 CTCCCTGCCATCCTTGGTGCCGTCCTGCCCCTTCCTCCTTCATGCCTGGGCCTGCAGGCC 2280
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Db      2221 CTCCCTGCCATCCTTGGTGCCGTCCTGCCCCTTCCTCCTTCATGCCTGGGCCTGCAGGCC 2280

Qy      2281 CACCCCAGCCACCACTGAGTCCACTCGGAGTGCCCTGTGTTCCTGGAGAAGGCATTCCAG 2340
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db      2281 CACCCCAGCCACCACTGAGTCCACTCGGAGTGCCCTGTGTTCCTGGAGAAGGCATTCCAG 2340

Qy      2341 GGTTGAATCTTGTCCCAGCCTCAGCCTGGGACACCTAGGTGGAGAGAGTGGTCTCCGCTC 2400
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Db      2341 GGTTGAATCTTGTCCCAGCCTCAGCCTGGGACACCTAGGTGGAGAGAGTGGTCTCCGCTC 2400

Qy      2401 TGAATTGGATCCAGGGGACCTGGGCTCATTCTTCTTGGCTCACCAACCCTGCAGGCCTCA 2460
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db      2401 TGAATTGGATCCAGGGGACCTGGGCTCATTCTTCTTGGCTCACCAACCCTGCAGGCCTCA 2460

Qy      2461 TCTTTCCCAAAACCCACTTTGTCTTGGTGGGAGTGGGTCCGCGCTGCTCTGCAGCAGGGG 2520
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db      2461 TCTTTCCCAAAACCCACTTTGTCTTGGTGGGAGTGGGTCCGCGCTGCTCTGCAGCAGGGG 2520

Qy      2521 CTGGGGAGTGGACAGCATCAGGTGGGAAAGTGGAGTCCACCCTCATGTTTCTGTAGGATT 2580
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db      2521 CTGGGGAGTGGACAGCATCAGGTGGGAAAGTGGAGTCCACCCTCATGTTTCTGTAGGATT 2580

Qy      2581 CTCACCGTGGGGCTGGAAGAAAAGAGCATCGACTTGATTTCTCCAACCACTCATCCCTCT 2640
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db      2581 CTCACCGTGGGGCTGGAAGAAAAGAGCATCGACTTGATTTCTCCAACCACTCATCCCTCT 2640

Qy      2641 TTTTCTTTCTTCCACCACTCCCCACCCCAGCTGTAGTTAATTTCAGTGCCTTACAAATCC 2700
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db      2641 TTTTCTTTCTTCCACCACTCCCCACCCCAGCTGTAGTTAATTTCAGTGCCTTACAAATCC 2700

Qy      2701 TAAGCTCAGAGAAAGTTCCATTTCCGTTCCAGAGGGAAGGGAACCTCCCTAGGTCCTTCC 2760
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Db      2701 TAAGCTCAGAGAAAGTTCCATTTCCGTTCCAGAGGGAAGGGAACCTCCCTAGGTCCTTCC 2760

Qy      2761 CTGGCTTGTTATAACGCAAAGCTTGGTTGTTTATGCAACTCTATCTTAAGAACTGCCCAG 2820
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Db      2761 CTGGCTTGTTATAACGCAAAGCTTGGTTGTTTATGCAACTCTATCTTAAGAACTGCCCAG 2820

Qy      2821 CCTCAGCTGAAAACCCGAATCTGAGAAGGAATTGCGTCATGTAAGGGAAGCTGGAATTAA 2880
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Db      2821 CCTCAGCTGAAAACCCGAATCTGAGAAGGAATTGCGTCATGTAAGGGAAGCTGGAATTAA 2880

Qy      2881 GGGAGCTGAGCCAGTCATGGTTGTGGCGTGTGAGTCAGGAGACCTAGGTTTCAGCCCCTC 2940
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db      2881 GGGAGCTGAGCCAGTCATGGTTGTGGCGTGTGAGTCAGGAGACCTAGGTTTCAGCCCCTC 2940

Qy      2941 TCTACTGTCAGCGAGCTGTGCAACGTGGGCAAGTCATTGTCCTCTGAGCTGCAGTTTCCT 3000
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Db      2941 TCTACTGTCAGCGAGCTGTGCAACGTGGGCAAGTCATTGTCCTCTGAGCTGCAGTTTCCT 3000

Qy      3001 CATCTGTCACATCGCTACAGACAAGACCTCCCTGGAACCCTTCTGATTGTCTTAGACACT 3060
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Db      3001 CATCTGTCACATCGCTACAGACAAGACCTCCCTGGAACCCTTCTGATTGTCTTAGACACT 3060

Qy      3061 GTGGTTGCAAAACCCACGGAAAGCCTCATTTGTGTGGAAAGTCAGAGGAAAAATGATCCA 3120
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Db      3061 GTGGTTGCAAAACCCACGGAAAGCCTCATTTGTGTGGAAAGTCAGAGGAAAAATGATCCA 3120

Qy      3121 GTGGACACTTGGGGATTATCTGTCATTCAAGATCCTTCCTTCAACCCCAAGGCCAGCTCC 3180
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Db      3121 GTGGACACTTGGGGATTATCTGTCATTCAAGATCCTTCCTTCAACCCCAAGGCCAGCTCC 3180

Qy      3181 CATCTCATTTCCAGAAAGGCTCATACCTGGCTTGCAGGGAAGCATCTGTCTTGTCATTCC 3240
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db      3181 CATCTCATTTCCAGAAAGGCTCATACCTGGCTTGCAGGGAAGCATCTGTCTTGTCATTCC 3240

Qy      3241 AGGTGCCAGAATCCTCTCAGAGTCATTGAAGGGTGTTCACCCATCCCACCCAAGGCTTGG 3300
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db      3241 AGGTGCCAGAATCCTCTCAGAGTCATTGAAGGGTGTTCACCCATCCCACCCAAGGCTTGG 3300

Qy      3301 CACACTGCCAGTGTCTTAGCAGGGTCTTGTGAGGGCTGGGGGCATCCAGGCACTCAGAAG 3360
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Db       3601 GGCCAGAGATGCCAGCAGCCTCTGTCGCCCTTAGTCATATAACCAAAATCCAGACCTTAT 3660

Qy       3661 CCACAACCCGGGGCTTGGAAAGGAAGGTATTTTGGAATCACACCCTCCGGTTATGTTGCT 3720
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       3661 CCACAACCCGGGGCTTGGAAAGGAAGGTATTTTGGAATCACACCCTCCGGTTATGTTGCT 3720

Qy       3721 CCAGTAAAATCTTGCCTGGAAAGAGGCAGTCTTCTTAGCATGGTGAGCTGAGTTCATGGC 3780
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       3721 CCAGTAAAATCTTGCCTGGAAAGAGGCAGTCTTCTTAGCATGGTGAGCTGAGTTCATGGC 3780

Qy       3781 TTTTTTTTGTAGCCAGTCCTGTCCCTGGCCATCCATGTGATGGTTTTGGATGGAGTTAAA 3840
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       3781 TTTTTTTTGTAGCCAGTCCTGTCCCTGGCCATCCATGTGATGGTTTTGGATGGAGTTAAA 3840

Qy       3841 CTTGATGCCAGTGGGCAGTGCATGTGGAAAGTATCAGAGTAAGCCTCTCCCCTCCAGAGC 3900
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       3841 CTTGATGCCAGTGGGCAGTGCATGTGGAAAGTATCAGAGTAAGCCTCTCCCCTCCAGAGC 3900

Qy       3901 CCTGAGTTTCTTGGCTGCATGAAGGTTTTCTTTAGAATCAGAATTGTAGCCAGTTTCTTT 3960
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       3901 CCTGAGTTTCTTGGCTGCATGAAGGTTTTCTTTAGAATCAGAATTGTAGCCAGTTTCTTT 3960

Qy       3961 GGCCAGAAGGATGAATACTTGGATATTACTGAAAGGGAGGGGTGGAGATGGGTGTGGCAG 4020
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       3961 GGCCAGAAGGATGAATACTTGGATATTACTGAAAGGGAGGGGTGGAGATGGGTGTGGCAG 4020

Qy       4021 TGTATGGTGTGTGATTTTTATTTTCTTCTTTGGTCATGGGGGCCAAGGAGAAAGGCATGA 4080
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       4021 TGTATGGTGTGTGATTTTTATTTTCTTCTTTGGTCATGGGGGCCAAGGAGAAAGGCATGA 4080

Qy       4081 ATCTTCCCTGTCAGGCTCTTACAGCCACAGGCACTGTGTCTACTGTCTGGAAGACATGTC 4140
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Db       4081 ATCTTCCCTGTCAGGCTCTTACAGCCACAGGCACTGTGTCTACTGTCTGGAAGACATGTC 4140

Qy       4141 CCCGTGGCTGTGGGGCCGCTGCTTCTGTTTAAATAAAAGTGGCCTGG 4187
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Db       4141 CCCGTGGCTGTGGGGCCGCTGCTTCTGTTTAAATAAAAGTGGCCTGG 4187


RESULT 4
US-09-873-319-290
; Sequence 290, Application US/09873319A
; Publication No. US20030134324A1
; GENERAL INFORMATION:
;  APPLICANT: Munger, William E.
;  APPLICANT: Kulkarni, Prakash
;  APPLICANT: Getzenberg, Robert H.
;  APPLICANT: Waga, Iwao
;  APPLICANT: Yamamoto, Jun
;  TITLE OF INVENTION: Identifying Drugs for and Diagnosis of Benign Prostatic
;  TITLE OF INVENTION: Hyperplasia Using Gene Expression Profiles
;  FILE REFERENCE: 44921-5029-US
;  CURRENT APPLICATION NUMBER: US/09/873,319A
;  CURRENT FILING DATE: 2001-06-05
;  EARLIER APPLICATION NUMBER: US 60/223,323
;  EARLIER FILING DATE: 2000-08-07
;  NUMBER OF SEQ ID NOS: 755
;  SOFTWARE: PatentIn Ver. 2.1
; SEQ ID NO 290
;   LENGTH: 4187
;   TYPE: DNA
;   ORGANISM: Homo sapiens
;   FEATURE:
;   OTHER INFORMATION: Genbank Accession No. US20030134324A1 D13643
US-09-873-319-290

  Query Match             100.0%;  Score 4187;  DB 10;  Length 4187;
  Best Local Similarity   100.0%;  Pred. No. 0;
  Matches 4187;  Conservative    0;  Mismatches    0;  Indels    0;  Gaps    0;

Qy          1 GGCGCGAACCCGCAGCGCTTACCGCGCGGCGCCGCACCATGGAGCCCGCCGTGTCGCTGG 60
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Db          1 GGCGCGAACCCGCAGCGCTTACCGCGCGGCGCCGCACCATGGAGCCCGCCGTGTCGCTGG 60

Qy         61 CCGTGTGCGCGCTGCTCTTCCTGCTGTGGGTGCGCCTGAAGGGGCTGGAGTTCGTGCTCA 120
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db         61 CCGTGTGCGCGCTGCTCTTCCTGCTGTGGGTGCGCCTGAAGGGGCTGGAGTTCGTGCTCA 120

Qy        121 TCCACCAGCGCTGGGTGTTCGTGTGCCTCTTCCTCCTGCCGCTCTCGCTTATCTTCGATA 180
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        121 TCCACCAGCGCTGGGTGTTCGTGTGCCTCTTCCTCCTGCCGCTCTCGCTTATCTTCGATA 180

Qy        181 TCTACTACTACGTGCGCGCCTGGGTGGTGTTCAAGCTCAGCAGCGCTCCGCGCCTGCACG 240
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        181 TCTACTACTACGTGCGCGCCTGGGTGGTGTTCAAGCTCAGCAGCGCTCCGCGCCTGCACG 240

Qy        241 AGCAGCGCGTGCGGGACATCCAGAAGCAGGTGCGGGAATGGAAGGAGCAGGGTAGCAAGA 300
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        241 AGCAGCGCGTGCGGGACATCCAGAAGCAGGTGCGGGAATGGAAGGAGCAGGGTAGCAAGA 300

Qy        301 CCTTCATGTGCACGGGGCGCCCTGGCTGGCTCACTGTCTCACTACGTGTCGGGAAGTACA 360
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        301 CCTTCATGTGCACGGGGCGCCCTGGCTGGCTCACTGTCTCACTACGTGTCGGGAAGTACA 360

Qy        361 AGAAGACACACAAAAACATCATGATCAACCTGATGGACATTCTGGAAGTGGACACCAAGA 420
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        361 AGAAGACACACAAAAACATCATGATCAACCTGATGGACATTCTGGAAGTGGACACCAAGA 420

Qy        421 AACAGATTGTCCGTGTGGAGCCCTTGGTGACCATGGGCCAGGTGACTGCCCTGCTGACCT 480
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        421 AACAGATTGTCCGTGTGGAGCCCTTGGTGACCATGGGCCAGGTGACTGCCCTGCTGACCT 480

Qy        481 CCATTGGCTGGACTCTCCCCGTGTTGCCTGAGCTTGATGACCTCACAGTGGGGGGCTTGA 540
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        481 CCATTGGCTGGACTCTCCCCGTGTTGCCTGAGCTTGATGACCTCACAGTGGGGGGCTTGA 540

Qy        541 TCATGGGCACAGGCATCGAGTCATCATCCCACAAGTACGGCCTGTTCCAACACATCTGCA 600
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        541 TCATGGGCACAGGCATCGAGTCATCATCCCACAAGTACGGCCTGTTCCAACACATCTGCA 600

Qy        601 CTGCTTACGAGCTGGTCCTGGCTGATGGCAGCTTTGTGCGATGCACTCCGTCCGAAAACT 660
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        601 CTGCTTACGAGCTGGTCCTGGCTGATGGCAGCTTTGTGCGATGCACTCCGTCCGAAAACT 660

Qy        661 CAGACCTGTTCTATGCCGTACCCTGGTCCTGTGGGACGCTGGGTTTCCTGGTGGCCGCTG 720
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Db        661 CAGACCTGTTCTATGCCGTACCCTGGTCCTGTGGGACGCTGGGTTTCCTGGTGGCCGCTG 720

Qy        721 AGATCCGCATCATCCCTGCCAAGAAGTACGTCAAGCTGCGTTTCGAGCCAGTGCGGGGCC 780
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        721 AGATCCGCATCATCCCTGCCAAGAAGTACGTCAAGCTGCGTTTCGAGCCAGTGCGGGGCC 780

Qy        781 TGGAGGCTATCTGTGCCAAGTTCACCCACGAGTCCCAGCGGCAGGAGAACCACTTCGTGG 840
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        781 TGGAGGCTATCTGTGCCAAGTTCACCCACGAGTCCCAGCGGCAGGAGAACCACTTCGTGG 840

Qy        841 AAGGGCTGCTCTACTCCCTGGATGAGGCTGTCATTATGACAGGGGTCATGACAGATGAGG 900
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        841 AAGGGCTGCTCTACTCCCTGGATGAGGCTGTCATTATGACAGGGGTCATGACAGATGAGG 900

Qy        901 CAGAGCCCAGCAAGCTGAATAGCATTGGCAATTACTACAAGCCGTGGTTCTTTAAGCATG 960
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Db        901 CAGAGCCCAGCAAGCTGAATAGCATTGGCAATTACTACAAGCCGTGGTTCTTTAAGCATG 960

Qy        961 TGGAGAACTATCTGAAGACAAACCGAGAGGGCCTGGAGTACATTCCCTTGAGACACTACT 1020
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        961 TGGAGAACTATCTGAAGACAAACCGAGAGGGCCTGGAGTACATTCCCTTGAGACACTACT 1020

Qy       1021 ACCACCGCCACACGCGCAGCATCTTCTGGGAGCTCCAGGACATCATCCCCTTTGGCAACA 1080
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Db       1021 ACCACCGCCACACGCGCAGCATCTTCTGGGAGCTCCAGGACATCATCCCCTTTGGCAACA 1080

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Qy       1441 AGATGCTGTATGCCGACTGCTACATGAACCGGGAGGAGTTCTGGGAGATGTTTGATGGCT 1500
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Qy       1501 CCTTGTACCACAAGCTGCGAGAGAAGCTGGGTTGCCAGGACGCCTTCCCCGAGGTGTACG 1560
Db       1501 CCTTGTACCACAAGCTGCGAGAGAAGCTGGGTTGCCAGGACGCCTTCCCCGAGGTGTACG 1560

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Db       1561 ACAAGATCTGCAAGGCCGCCAGGCACTGAGCTGGAGCCCGCCTGGAGAGACAGACACGTG 1620

Qy       1621 TGAGTGGTCAGGCATCTTCCCTTCACTCAAGCTTGGCTGCTTTCCTAGATCCACACTTTC 1680
Db       1621 TGAGTGGTCAGGCATCTTCCCTTCACTCAAGCTTGGCTGCTTTCCTAGATCCACACTTTC 1680

Qy       1681 AAAGAGAAACCCCTCCAGAACTCCCACCCTGACAGCCCAACACCACCTTCCTCCTGGCTT 1740
Db       1681 AAAGAGAAACCCCTCCAGAACTCCCACCCTGACAGCCCAACACCACCTTCCTCCTGGCTT 1740

Qy       1741 CCAGGGGGCAGCCCAGTGGAATGGAAAGAATGTGGGATTTGGAGTCAGACAAGCCTGAGT 1800
Db       1741 CCAGGGGGCAGCCCAGTGGAATGGAAAGAATGTGGGATTTGGAGTCAGACAAGCCTGAGT 1800

Qy       1801 CCAGTTCCCCGTTTAGAACTCATTAGCTGTGTGACTCTGGGTGAGTCCCTTAACCCCTCT 1860
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Qy       1861 GAGCCCGGGTCTCTTCATTAGTTGAAAGGGATAGTAATACCTACTTGCAGGTTGTTGTCA 1920
Db       1861 GAGCCCGGGTCTCTTCATTAGTTGAAAGGGATAGTAATACCTACTTGCAGGTTGTTGTCA 1920

Qy       1921 TCTGAGTTGAGCACTGGTCACATTGAAGGTGCTGGGTAAGTGGTAGCTCTTGTTGCTTCC 1980
Db       1921 TCTGAGTTGAGCACTGGTCACATTGAAGGTGCTGGGTAAGTGGTAGCTCTTGTTGCTTCC 1980

Qy       1981 CGTTCAGCGTCACATCTGCAGTGGAGCCTGAAAAGGCTCCACATTAGGTCACCTGTGCAC 2040
Db       1981 CGTTCAGCGTCACATCTGCAGTGGAGCCTGAAAAGGCTCCACATTAGGTCACCTGTGCAC 2040

Qy       2041 AGCCATGGCTGGAATGATGAAGGGGATACGCTGGAGTTGCCCTGCCATCGCCTCCATCAG 2100
Db       2041 AGCCATGGCTGGAATGATGAAGGGGATACGCTGGAGTTGCCCTGCCATCGCCTCCATCAG 2100

Qy       2101 CCAGACGAGGTCCTCACAGGAGAAGGACAGCTCTTCCCCACCCTGGGATCTCAGGAGGGC 2160
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Qy       2161 AGCCACGGAGTGGGGAGGCCCCAGATGCGCTGTGCCAAAGCCAGGTCCGAGGCCAAAGTT 2220
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Qy       2221 CTCCCTGCCATCCTTGGTGCCGTCCTGCCCCTTCCTCCTTCATGCCTGGGCCTGCAGGCC 2280
Db       2221 CTCCCTGCCATCCTTGGTGCCGTCCTGCCCCTTCCTCCTTCATGCCTGGGCCTGCAGGCC 2280

Qy       2281 CACCCCAGCCACCACTGAGTCCACTCGGAGTGCCCTGTGTTCCTGGAGAAGGCATTCCAG 2340
Db       2281 CACCCCAGCCACCACTGAGTCCACTCGGAGTGCCCTGTGTTCCTGGAGAAGGCATTCCAG 2340

Qy       2341 GGTTGAATCTTGTCCCAGCCTCAGCCTGGGACACCTAGGTGGAGAGAGTGGTCTCCGCTC 2400
Db       2341 GGTTGAATCTTGTCCCAGCCTCAGCCTGGGACACCTAGGTGGAGAGAGTGGTCTCCGCTC 2400

Qy       2401 TGAATTGGATCCAGGGGACCTGGGCTCATTCTTCTTGGCTCACCAACCCTGCAGGCCTCA 2460
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Qy       2461 TCTTTCCCAAAACCCACTTTGTCTTGGTGGGAGTGGGTCCGCGCTGCTCTGCAGCAGGGG 2520
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Qy       2521 CTGGGGAGTGGACAGCATCAGGTGGGAAAGTGGAGTCCACCCTCATGTTTCTGTAGGATT 2580
Db       2521 CTGGGGAGTGGACAGCATCAGGTGGGAAAGTGGAGTCCACCCTCATGTTTCTGTAGGATT 2580

Qy       2581 CTCACCGTGGGGCTGGAAGAAAAGAGCATCGACTTGATTTCTCCAACCACTCATCCCTCT 2640
Db       2581 CTCACCGTGGGGCTGGAAGAAAAGAGCATCGACTTGATTTCTCCAACCACTCATCCCTCT 2640

Qy       2641 TTTTCTTTCTTCCACCACTCCCCACCCCAGCTGTAGTTAATTTCAGTGCCTTACAAATCC 2700
Db       2641 TTTTCTTTCTTCCACCACTCCCCACCCCAGCTGTAGTTAATTTCAGTGCCTTACAAATCC 2700

Qy       2701 TAAGCTCAGAGAAAGTTCCATTTCCGTTCCAGAGGGAAGGGAACCTCCCTAGGTCCTTCC 2760
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Qy       2761 CTGGCTTGTTATAACGCAAAGCTTGGTTGTTTATGCAACTCTATCTTAAGAACTGCCCAG 2820
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Qy       2821 CCTCAGCTGAAAACCCGAATCTGAGAAGGAATTGCGTCATGTAAGGGAAGCTGGAATTAA 2880
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Qy       2881 GGGAGCTGAGCCAGTCATGGTTGTGGCGTGTGAGTCAGGAGACCTAGGTTTCAGCCCCTC 2940
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Qy       3001 CATCTGTCACATCGCTACAGACAAGACCTCCCTGGAACCCTTCTGATTGTCTTAGACACT 3060
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Qy       3061 GTGGTTGCAAAACCCACGGAAAGCCTCATTTGTGTGGAAAGTCAGAGGAAAAATGATCCA 3120
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Qy       3121 GTGGACACTTGGGGATTATCTGTCATTCAAGATCCTTCCTTCAACCCCAAGGCCAGCTCC 3180
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Qy       3181 CATCTCATTTCCAGAAAGGCTCATACCTGGCTTGCAGGGAAGCATCTGTCTTGTCATTCC 3240
Db       3181 CATCTCATTTCCAGAAAGGCTCATACCTGGCTTGCAGGGAAGCATCTGTCTTGTCATTCC 3240

Qy       3241 AGGTGCCAGAATCCTCTCAGAGTCATTGAAGGGTGTTCACCCATCCCACCCAAGGCTTGG 3300
Db       3241 AGGTGCCAGAATCCTCTCAGAGTCATTGAAGGGTGTTCACCCATCCCACCCAAGGCTTGG 3300

Qy       3301 CACACTGCCAGTGTCTTAGCAGGGTCTTGTGAGGGCTGGGGGCATCCAGGCACTCAGAAG 3360
Db       3301 CACACTGCCAGTGTCTTAGCAGGGTCTTGTGAGGGCTGGGGGCATCCAGGCACTCAGAAG 3360

Qy       3361 GCAAAGGAACCACCCTACCCATTTGGCCTCTGGAGGGGGCAGAAGAAAGAAAGAAACCTC 3420
Db       3361 GCAAAGGAACCACCCTACCCATTTGGCCTCTGGAGGGGGCAGAAGAAAGAAAGAAACCTC 3420

Qy       3421 ATCCTATATTTTACAAAGCATGTGAATTCTGGCATTAGCTCTCATAGGAGACCCATGTGC 3480
Db       3421 ATCCTATATTTTACAAAGCATGTGAATTCTGGCATTAGCTCTCATAGGAGACCCATGTGC 3480

Qy       3481 TTCCTTGCTCAGTGCAAAACTGATGATTCTACTTGCTGTAGATGAATGGTTAACACGAGC 3540
Db       3481 TTCCTTGCTCAGTGCAAAACTGATGATTCTACTTGCTGTAGATGAATGGTTAACACGAGC 3540

Qy       3541 TAGTTAAACAGTGCCATTGTTTTGCCAGTGAAGCCTCCAACCCTAAGCCACTGGGACGGT 3600
Db       3541 TAGTTAAACAGTGCCATTGTTTTGCCAGTGAAGCCTCCAACCCTAAGCCACTGGGACGGT 3600

Qy       3601 GGCCAGAGATGCCAGCAGCCTCTGTCGCCCTTAGTCATATAACCAAAATCCAGACCTTAT 3660
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Qy       3901 CCTGAGTTTCTTGGCTGCATGAAGGTTTTCTTTAGAATCAGAATTGTAGCCAGTTTCTTT 3960
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Db       3901 CCTGAGTTTCTTGGCTGCATGAAGGTTTTCTTTAGAATCAGAATTGTAGCCAGTTTCTTT 3960

Qy       3961 GGCCAGAAGGATGAATACTTGGATATTACTGAAAGGGAGGGGTGGAGATGGGTGTGGCAG 4020
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       3961 GGCCAGAAGGATGAATACTTGGATATTACTGAAAGGGAGGGGTGGAGATGGGTGTGGCAG 4020

Qy       4021 TGTATGGTGTGTGATTTTTATTTTCTTCTTTGGTCATGGGGGCCAAGGAGAAAGGCATGA 4080
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       4021 TGTATGGTGTGTGATTTTTATTTTCTTCTTTGGTCATGGGGGCCAAGGAGAAAGGCATGA 4080

Qy       4081 ATCTTCCCTGTCAGGCTCTTACAGCCACAGGCACTGTGTCTACTGTCTGGAAGACATGTC 4140
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Db       4081 ATCTTCCCTGTCAGGCTCTTACAGCCACAGGCACTGTGTCTACTGTCTGGAAGACATGTC 4140

Qy       4141 CCCGTGGCTGTGGGGCCGCTGCTTCTGTTTAAATAAAAGTGGCCTGG 4187
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Db       4141 CCCGTGGCTGTGGGGCCGCTGCTTCTGTTTAAATAAAAGTGGCCTGG 4187


RESULT 3
US-09-960-706-477
; Sequence 477, Application US/09960706
; Publication No. US20030134280A1
; GENERAL INFORMATION:
;  APPLICANT: Munger, William E.
;  TITLE OF INVENTION: Identifying Drugs for and Diagnosis of Benign Prostatic Hyperplas
;  TITLE OF INVENTION:  Gene Expression Profiles
;  FILE REFERENCE: 44921-5029-01US
;  CURRENT APPLICATION NUMBER: US/09/960,706
;  CURRENT FILING DATE:  2001-09-24
;  PRIOR APPLICATION NUMBER: 60/223,323
;  PRIOR FILING DATE: 2000-08-07
;  PRIOR APPLICATION NUMBER: 09/873,319
;  PRIOR FILING DATE: 2001-06-05
;  NUMBER OF SEQ ID NOS: 1124
;  SOFTWARE: PatentIn Ver. 2.1
; SEQ ID NO 477
;   LENGTH: 4187
;   TYPE: DNA
;   ORGANISM: Homo sapiens
;   FEATURE:
;   OTHER INFORMATION: Genbank Accession No. US20030134280A1 D13643
US-09-960-706-477

  Query Match             100.0%;  Score 4187;  DB 10;  Length 4187;
  Best Local Similarity   100.0%;  Pred. No. 0;
  Matches 4187;  Conservative    0;  Mismatches    0;  Indels    0;  Gaps    0;

Qy          1 GGCGCGAACCCGCAGCGCTTACCGCGCGGCGCCGCACCATGGAGCCCGCCGTGTCGCTGG 60
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Qy         61 CCGTGTGCGCGCTGCTCTTCCTGCTGTGGGTGCGCCTGAAGGGGCTGGAGTTCGTGCTCA 120
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Qy        121 TCCACCAGCGCTGGGTGTTCGTGTGCCTCTTCCTCCTGCCGCTCTCGCTTATCTTCGATA 180
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Qy        181 TCTACTACTACGTGCGCGCCTGGGTGGTGTTCAAGCTCAGCAGCGCTCCGCGCCTGCACG 240
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Qy        241 AGCAGCGCGTGCGGGACATCCAGAAGCAGGTGCGGGAATGGAAGGAGCAGGGTAGCAAGA 300
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Db        241 AGCAGCGCGTGCGGGACATCCAGAAGCAGGTGCGGGAATGGAAGGAGCAGGGTAGCAAGA 300

Qy        301 CCTTCATGTGCACGGGGCGCCCTGGCTGGCTCACTGTCTCACTACGTGTCGGGAAGTACA 360
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Qy        481 CCATTGGCTGGACTCTCCCCGTGTTGCCTGAGCTTGATGACCTCACAGTGGGGGGCTTGA 540
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Qy        601 CTGCTTACGAGCTGGTCCTGGCTGATGGCAGCTTTGTGCGATGCACTCCGTCCGAAAACT 660
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Db        601 CTGCTTACGAGCTGGTCCTGGCTGATGGCAGCTTTGTGCGATGCACTCCGTCCGAAAACT 660

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Db        901 CAGAGCCCAGCAAGCTGAATAGCATTGGCAATTACTACAAGCCGTGGTTCTTTAAGCATG 960

Qy        961 TGGAGAACTATCTGAAGACAAACCGAGAGGGCCTGGAGTACATTCCCTTGAGACACTACT 1020
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Qy       1021 ACCACCGCCACACGCGCAGCATCTTCTGGGAGCTCCAGGACATCATCCCCTTTGGCAACA 1080
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Db       1021 ACCACCGCCACACGCGCAGCATCTTCTGGGAGCTCCAGGACATCATCCCCTTTGGCAACA 1080

Qy       1081 ACCCCATCTTCCGCTACCTCTTTGGCTGGATGGTGCCTCCCAAGATCTCCCTCCTGAAGC 1140
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Qy       1141 TGACCCAGGGTGAGACCCTGCGCAAGCTGTACGAGCAGCACCACGTGGTGCAGGACATGC 1200
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Db       1141 TGACCCAGGGTGAGACCCTGCGCAAGCTGTACGAGCAGCACCACGTGGTGCAGGACATGC 1200

Qy       1201 TGGTGCCCATGAAGTGCCTGCAGCAGGCCCTGCACACCTTCCAAAACGACATCCACGTCT 1260
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Db       1201 TGGTGCCCATGAAGTGCCTGCAGCAGGCCCTGCACACCTTCCAAAACGACATCCACGTCT 1260

Qy       1261 ACCCCATCTGGCTGTGTCCGTTCATCCTGCCCAGCCAGCCAGGCCTAGTGCACCCCAAAG 1320
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Db       1261 ACCCCATCTGGCTGTGTCCGTTCATCCTGCCCAGCCAGCCAGGCCTAGTGCACCCCAAAG 1320

Qy       1321 GAAATGAGGCAGAGCTCTACATCGACATTGGAGCATATGGGGAGCCGCGTGTGAAACACT 1380
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       1321 GAAATGAGGCAGAGCTCTACATCGACATTGGAGCATATGGGGAGCCGCGTGTGAAACACT 1380

Qy       1381 TTGAAGCCAGGTCCTGCATGAGGCAGCTGGAGAAGTTTGTCCGCAGCGTGCATGGCTTCC 1440
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Db       1381 TTGAAGCCAGGTCCTGCATGAGGCAGCTGGAGAAGTTTGTCCGCAGCGTGCATGGCTTCC 1440
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Db      1681 AAAGAGAAACCCCTCCAGAACTCCCACCCTGACAGCCCAACACCACCTTCCTCCTGGCTT 1740
Qy      1741 CCAGGGGGCAGCCCAGTGGAATGGAAAGAATGTGGGATTTGGAGTCAGACAAGCCTGAGT 1800
Db      1741 CCAGGGGGCAGCCCAGTGGAATGGAAAGAATGTGGGATTTGGAGTCAGACAAGCCTGAGT 1800
Qy      1801 CCAGTTCCCCGTTTAGAACTCATTAGCTGTGTGACTCTGGGTGAGTCCCTTAACCCCTCT 1860
Db      1801 CCAGTTCCCCGTTTAGAACTCATTAGCTGTGTGACTCTGGGTGAGTCCCTTAACCCCTCT 1860
Qy      1861 GAGCCCGGGTCTCTTCATTAGTTGAAAGGGATAGTAATACCTACTTGCAGGTTGTTGTCA 1920
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Qy      1921 TCTGAGTTGAGCACTGGTCACATTGAAGGTGCTGGGTAAGTGGTAGCTCTTGTTGCTTCC 1980
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Qy      1981 CGTTCAGCGTCACATCTGCAGTGGAGCCTGAAAAGGCTCCACATTAGGTCACCTGTGCAC 2040
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Qy      2041 AGCCATGGCTGGAATGATGAAGGGGATACGCTGGAGTTGCCCTGCCATCGCCTCCATCAG 2100
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Qy      2161 AGCCACGGAGTGGGGAGGCCCCAGATGCGCTGTGCCAAAGCCAGGTCCGAGGCCAAAGTT 2220
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Qy      2341 GGTTGAATCTTGTCCCAGCCTCAGCCTGGGACACCTAGGTGGAGAGAGTGGTCTCCGCTC 2400
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Db      2521 CTGGGGAGTGGACAGCATCAGGTGGGAAAGTGGAGTCCACCCTCATGTTTCTGTAGGATT 2580
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Qy      3181 CATCTCATTTCCAGAAAGGCTCATACCTGGCTTGCAGGGAAGCATCTGTCTTGTCATTCC 3240
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RESULT 2
US-09-880-107-1609
; Sequence 1609, Application US/09880107
; Patent No. US20020142981A1
; GENERAL INFORMATION:
; APPLICANT: Horne, Darci T.
; APPLICANT: Vockley, Joseph G.
; APPLICANT: Scherf, Uwe
; APPLICANT: Gene Logic, Inc.
; TITLE OF INVENTION: Gene Expression Profiles in Liver Cancer
; FILE REFERENCE: 44921-5028-WO
; CURRENT APPLICATION NUMBER: US/09/880,107
; CURRENT FILING DATE: 2001-06-14
; PRIOR APPLICATION NUMBER: US 60/211,379
; PRIOR FILING DATE: 2000-06-14
; PRIOR APPLICATION NUMBER: US 60/237,054
; PRIOR FILING DATE: 2000-10-02
; NUMBER OF SEQ ID NOS: 3950
; SOFTWARE: PatentIn Ver. 2.1
; SEQ ID NO 1609
; LENGTH: 4187
; TYPE: DNA
; ORGANISM: Homo sapiens
; FEATURE:
; OTHER INFORMATION: Genbank Accession No. US20020142981A1 D13643
US-09-880-107-1609

Query Match 100.0%; Score 4187; DB 9; Length 4187;
Best Local Similarity 100.0%; Pred. No. 0;
Matches 4187; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

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Qy        781 TGGAGGCTATCTGTGCCAAGTTCACCCACGAGTCCCAGCGGCAGGAGAACCACTTCGTGG 840
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Db      4141 CCCGTGGCTGTGGGGCCGCTGCTTCTGTTTAAATAAAAGTGGCCTGG 4187
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;  PRIOR FILING DATE: 2000-09-27
;  NUMBER OF SEQ ID NOS: 2276
;  SOFTWARE: PatentIn version 3.0
; SEQ ID NO 2112
;   LENGTH: 4187
;   TYPE: DNA
;   ORGANISM: Homo sapiens
US-09-954-456-2112

  Query Match            100.0%;  Score 4187;  DB 9;  Length 4187;
  Best Local Similarity  100.0%;  Pred. No. 0;
  Matches 4187;  Conservative    0;  Mismatches    0;  Indels    0;  Gaps    0;

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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

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1	4187	100.0	4187	9	US-09-954-456-2112	Sequence 2112, Ap
2	4187	100.0	4187	9	US-09-880-107-1609	Sequence 1609, Ap
3	4187	100.0	4187	10	US-09-960-706-477	Sequence 477, App
4	4187	100.0	4187	10	US-09-873-319-290	Sequence 290, App
5	4187	100.0	4187	17	US-10-172-118-355	Sequence 355, App
6	4187	100.0	4187	18	US-10-240-425-1203	Sequence 1203, Ap
7	4187	100.0	4187	18	US-10-342-887-355	Sequence 355, App
8	4187	100.0	4187	21	US-10-843-641A-5139	Sequence 5139, Ap
9	4187	100.0	4187	22	US-10-934-998-53	Sequence 53, Appl
10	4187	100.0	4248	21	US-10-956-157-1950	Sequence 1950, Ap
11	4187	100.0	4248	22	US-10-934-998-32	Sequence 32, Appl
12	4183.8	99.9	4275	14	US-10-176-847-1	Sequence 1, Appli
13	4182.2	99.9	4248	13	US-10-087-192-1259	Sequence 1259, Ap
14	4067.8	97.2	4219	21	US-10-479-081-121	Sequence 121, App
15	4058.8	96.9	4094	21	US-10-857-942-3	Sequence 3, Appli
16	3874.4	92.5	4525	16	US-10-133-013-212	Sequence 212, App
17	2750.4	65.7	57564	13	US-10-087-192-1258	Sequence 1258, Ap
18	2577.6	61.6	2683	9	US-09-925-301-387	Sequence 387, App
19	2315.4	55.3	2339	19	US-10-398-037-80	Sequence 80, Appl
c 20	1774.6	42.4	1972	13	US-10-027-632-99214	Sequence 99214, A
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26	561.4	13.4	728	17	US-10-220-891-71	Sequence 71, Appl
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28	431.4	10.3	474	10	US-09-918-995-917	Sequence 917, App
29	406.2	9.7	497	17	US-10-242-535A-41864	Sequence 41864, A
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c 31	388.2	9.3	440	10	US-09-918-995-34917	Sequence 34917, A
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c 33	359	8.6	444	10	US-09-918-995-34919	Sequence 34919, A
c 34	303.4	7.2	463	18	US-10-240-425-815	Sequence 815, App
c 35	295.6	7.1	5758	22	US-10-934-998-179	Sequence 179, App
c 36	294.8	7.0	298	22	US-10-934-998-133	Sequence 133, App
37	291	7.0	2074	20	US-10-739-930-4489	Sequence 4489, Ap
38	283.2	6.8	46675	13	US-10-087-192-1255	Sequence 1255, Ap
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40	220.8	5.3	2179	19	US-10-767-701-15414	Sequence 15414, A
41	214	5.1	1683	18	US-10-395-463-19	Sequence 19, Appl
42	214	5.1	1686	17	US-10-259-194A-333	Sequence 333, App
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44	210.2	5.0	1689	17	US-10-259-194A-620	Sequence 620, App
c 45	205	4.9	271	9	US-09-864-761-18582	Sequence 18582, A

ALIGNMENTS

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; Sequence 2112, Application US/09954456
; Patent No. US20020115057A1
; GENERAL INFORMATION:
; APPLICANT: Young, Paul
; TITLE OF INVENTION: Process for Identifying Anti-Cancer Therapeutic Agents Using Can
; TITLE OF INVENTION: Sets
; FILE REFERENCE: 689290-76
; CURRENT APPLICATION NUMBER: US/09/954,456
; CURRENT FILING DATE: 2001-09-18
; PRIOR APPLICATION NUMBER: US/60/233,617
; PRIOR FILING DATE: 2000-09-18
; PRIOR APPLICATION NUMBER: US/60/234,052
; PRIOR FILING DATE: 2000-09-20
; PRIOR APPLICATION NUMBER: US/60/234,923
; PRIOR FILING DATE: 2000-09-25
; PRIOR APPLICATION NUMBER: US/60/235,134
; PRIOR FILING DATE: 2000-09-25
; PRIOR APPLICATION NUMBER: US/60/235,637
; PRIOR FILING DATE: 2000-09-26
; PRIOR APPLICATION NUMBER: US/60/235,638
; PRIOR FILING DATE: 2000-09-26
; PRIOR APPLICATION NUMBER: US/60/235,711
; PRIOR FILING DATE: 2000-09-27
; PRIOR APPLICATION NUMBER: US/60/235,720
; PRIOR FILING DATE: 2000-09-27
; PRIOR APPLICATION NUMBER: US/60/235,840
; PRIOR FILING DATE: 2000-09-27
; PRIOR APPLICATION NUMBER: US/60/235,863

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Db      19474 C 19474


RESULT 14
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; Sequence 20, Application US/09230371A
; Patent No. 6348586
; GENERAL INFORMATION:
;  APPLICANT:  Chang, Yuan
;  APPLICANT:  Bohenzky, Roy A
;  APPLICANT:  Russo, James J
;  APPLICANT:  Edelman, Isidore S
;  APPLICANT:  Moore, Patrick S
;  TITLE OF INVENTION:  UNIQUE ASSOCIATED KAPOSI'S SARCOMA VIRUS SEQUENCES AND
;  TITLE OF INVENTION:  USES THEREOF
;  FILE REFERENCE:  45185-G-PCT-US
;  CURRENT APPLICATION NUMBER: US/09/230,371A
;  CURRENT FILING DATE:  1999-11-17
;  PRIOR APPLICATION NUMBER:  PCT/US97/13346
;  PRIOR FILING DATE:  1997-07-22
;  NUMBER OF SEQ ID NOS:  30
;  SOFTWARE:  PatentIn Ver. 2.0
; SEQ ID NO 20
;   LENGTH: 32207
;   TYPE: DNA
;   ORGANISM: Kaposi's sarcoma-associated herpesvirus
US-09-230-371A-20

  Query Match             1.3%;  Score 54.6;  DB 3;  Length 32207;
  Best Local Similarity   56.4%;  Pred. No. 0.0022;
  Matches  102;  Conservative    0;  Mismatches   79;  Indels    0;  Gaps    0;

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Qy        252 C 252
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Db      19474 C 19474


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; Sequence 99, Application US/09573080A
; Patent No. 6828097
; GENERAL INFORMATION:
;  APPLICANT:  JOAN, KNOLL
;  APPLICANT:  ROGAN, PETER
;  TITLE OF INVENTION:  SINGLE COPY GENOMIC HYBRIDIZATION PROBES AND METHOD OF GENERATI
;  FILE REFERENCE:  30307
;  CURRENT APPLICATION NUMBER: US/09/573,080A
;  CURRENT FILING DATE:  2000-05-16
;  NUMBER OF SEQ ID NOS:  479
;  SOFTWARE:  PatentIn version 3.0
; SEQ ID NO 99
;   LENGTH: 262
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;   ORGANISM: Homo sapiens
;   FEATURE:
;   NAME/KEY: repeat_region
;   LOCATION: (1)..(262)
;   OTHER INFORMATION: mir
; PUBLICATION INFORMATION:
; PUBLICATION INFORMATION:
;   AUTHORS: Jurka, J; Walichiewicz, J; Milosavljevic, A
;   TITLE: Prototypic sequences for human repetitive DNA
;   JOURNAL: Journal of Molecular Evolution
;   VOLUME: 35
;   ISSUE: 4
;   PAGES: 286-291
;   DATE: 1992-10-__
;   DATABASE ACCESSION NUMBER: Database of repetitive elements (repbase)
;   DATABASE ENTRY DATE: ____-__-__
;   DATABASE ENTRY DATE: 1996-01-26
US-09-573-080A-99

  Query Match             1.3%;  Score 54.2;  DB 4;  Length 262;
  Best Local Similarity   64.0%;  Pred. No. 0.00016;
  Matches   80;  Conservative    1;  Mismatches   44;  Indels    0;  Gaps    0;

Qy       1766 AAGAATGTGGGATTTGGAGTCAGACAAGCCTGAGTCCAGTTCCCCGTTTAGAACTCATTA 1825
              ||||    ||: | ||||| |||||   |  |  || | | ||   | |   ||| | ||
Db         20 AAGAGCACGGRCTCTGGAGCCAGACTGCCTGGGTTCGAATCCCGGCTCTGCCACTTACTA 79

Qy       1826 GCTGTGTGACTCTGGGTGAGTCCCTTAACCCCTCTGAGCCCGGGTCTCTTCATTAGTTGA 1885
              ||||||||||  ||||  |||  ||||||| ||||| |||   || || ||||  ||  |
Db         80 GCTGTGTGACCTTGGGCAAGTTACTTAACCTCTCTGTGCCTCAGTTTCCTCATCTGTAAA 139

Qy       1886 AAGGG 1890
              | |||
Db        140 ATGGG 144


Search completed: August 17, 2005, 18:01:19
Job time : 639.158 secs
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;   ORGANISM: Kaposi's sarcoma-associated herpesvirus
US-09-894-273-1

  Query Match             1.3%;  Score 54.6;  DB 4;  Length 3489;
  Best Local Similarity   56.4%;  Pred. No. 0.00058;
  Matches  102;  Conservative    0;  Mismatches   79;  Indels    0;  Gaps    0;

Qy          72 CTGCTCTTCCTGCTGTGGGTGCGCCTGAAGGGGCTGGAGTTCGTGCTCATCCACCAGCGC 131
               ||||||||||||||| |      | |||  |     |||   ||| ||||| ||||||  | |
Db        2703 CTGCTCTTCCTGCTCTTCCACCTCCTCTAACTCCTGCTCTTCCTGCTCTTCCACCTCCTC 2644

Qy         132 TGGGTGTTCGTGTGCCTCTTCCTCCTGCCGCTCTCGCTTATCTTCGATATCTACTACTAC 191
               |   | | |   |||  |||||||||  ||      |  || || |  ||  ||| | |
Db        2643 TAACTCCTGCTCCTCCTGCTCCTCCTGCTCCTGCTCTTGCTCCTCCACCTCCTCTAATTC 2584

Qy         192 GTGCGCGCCTGGGTGGTGTTCAAGCTCAGCAGCGCTCCGCGCCTGCACGAGCAGCGCGTG 251
               ||| |  |  | | || ||  ||||  |  |   | || | ||| |   || | | ||
Db        2583 CTGCTCTTCCTGCTCCTGCTCTTGCTCTTCCACCTCCTGCTCTTGCTCTTCCACCTCCTG 2524

Qy         252 C 252
               |
Db        2523 C 2523


RESULT 12
US-08-770-379-20
; Sequence 20, Application US/08770379
; Patent No. 5849564
;  GENERAL INFORMATION:
;    APPLICANT:  Chang, Yuan
;    APPLICANT:  Bohenzky, Roy A.
;    APPLICANT:  Russo, James J.
;    APPLICANT:  Edelman, Isidore S.
;    APPLICANT:  Moore, Patrick S.
;    TITLE OF INVENTION:  POLYPEPTIDES FROM KAPOSI'S SARCOMA-ASSOCIATED
;    TITLE OF INVENTION:  HERPESVIRUS, DNA ENCODING SAME AND USES THEREOF
;    NUMBER OF SEQUENCES:  20
;    CORRESPONDENCE ADDRESS:
;      ADDRESSEE:  Cooper & Dunham LLP
;      STREET:  1185 Avenue of the Americas
;      CITY:  New York
;      STATE:  New York
;      COUNTRY:  U.S.A.
;      ZIP:  10036
;    COMPUTER READABLE FORM:
;      MEDIUM TYPE:  Floppy disk
;      COMPUTER:  IBM PC compatible
;      OPERATING SYSTEM:  PC-DOS/MS-DOS
;      SOFTWARE:  PatentIn Release #1.0, Version #1.30
;    CURRENT APPLICATION DATA:
;      APPLICATION NUMBER:  US/08/770,379
;      FILING DATE:
;      CLASSIFICATION:  435
;    ATTORNEY/AGENT INFORMATION:
;      NAME:  White, John P.
;      REGISTRATION NUMBER:  28,678
;      REFERENCE/DOCKET NUMBER:  52342
;    TELECOMMUNICATION INFORMATION:
;      TELEPHONE:  (212) 278-0400
;      TELEFAX:  (212) 391-0525
;  INFORMATION FOR SEQ ID NO:  20:
;    SEQUENCE CHARACTERISTICS:
;      LENGTH:  32207 base pairs
;      TYPE:  nucleic acid
;      STRANDEDNESS:  double
;      TOPOLOGY:  linear
;    MOLECULE TYPE:  DNA (genomic)
US-08-770-379-20

  Query Match             1.3%;  Score 54.6;  DB 2;  Length 32207;
  Best Local Similarity   56.4%;  Pred. No. 0.0022;
  Matches  102;  Conservative    0;  Mismatches   79;  Indels    0;  Gaps    0;

Qy          72 CTGCTCTTCCTGCTGTGGGTGCGCCTGAAGGGGCTGGAGTTCGTGCTCATCCACCAGCGC 131
               ||||||||||||||| |      | |||  |     |||   ||| ||||| ||||||  | |
Db       19294 CTGCTCTTCCTGCTCTTCCACCTCCTCTAACTCCTGCTCTTCCTGCTCTTCCACCTCCTC 19353

Qy         132 TGGGTGTTCGTGTGCCTCTTCCTCCTGCCGCTCTCGCTTATCTTCGATATCTACTACTAC 191
               |   | | |   |||  |||||||||  ||      |  || || |  ||  ||| | |
Db       19354 TAACTCCTGCTCCTCCTGCTCCTCCTGCTCCTGCTCTTGCTCCTCCACCTCCTCTAATTC 19413

Qy         192 GTGCGCGCCTGGGTGGTGTTCAAGCTCAGCAGCGCTCCGCGCCTGCACGAGCAGCGCGTG 251
               ||| |  |  | | || ||  ||||  |  |   | || | ||| |   || | | ||
Db       19414 CTGCTCTTCCTGCTCCTGCTCTTGCTCTTCCACCTCCTGCTCTTGCTCTTCCACCTCCTG 19473

Qy         252 C 252
               |
Db       19474 C 19474


RESULT 13
US-08-757-669A-20
; Sequence 20, Application US/08757669A
; Patent No. 6183751
;  GENERAL INFORMATION:
;    APPLICANT:  Chang, Yuan
;    APPLICANT:  Bohenzky, Roy A.
;    APPLICANT:  Russo, James J.
;    APPLICANT:  Edelman, Isidore S.
;    APPLICANT:  Moore, Patrick S.
;    TITLE OF INVENTION:  UNIQUE ASSOCIATED KAPOSI'S SARCOMA VIRUS
;    TITLE OF INVENTION:  SEQUENCES AND USES THEREOF
;    NUMBER OF SEQUENCES:  20
;    CORRESPONDENCE ADDRESS:
;      ADDRESSEE:  Cooper & Dunham LLP
;      STREET:  1185 Avenue of the Americas
;      CITY:  New York
;      STATE:  New York
;      COUNTRY:  U.S.A.
;      ZIP:  10036
;    COMPUTER READABLE FORM:
;      MEDIUM TYPE:  Floppy disk
;      COMPUTER:  IBM PC compatible
;      OPERATING SYSTEM:  PC-DOS/MS-DOS
;      SOFTWARE:  PatentIn Release #1.0, Version #1.30
;    CURRENT APPLICATION DATA:
;      APPLICATION NUMBER:  US/08/757,669A
;      FILING DATE:
;      CLASSIFICATION:  424
;    ATTORNEY/AGENT INFORMATION:
;      NAME:  White, John P.
;      REGISTRATION NUMBER:  28,678
;      REFERENCE/DOCKET NUMBER:  45185-F
;    TELECOMMUNICATION INFORMATION:
;      TELEPHONE:  (212) 278-0400
;      TELEFAX:  (212) 391-0525
;  INFORMATION FOR SEQ ID NO:  20:
;    SEQUENCE CHARACTERISTICS:
;      LENGTH:  32207 base pairs
;      TYPE:  nucleic acid
;      STRANDEDNESS:  double
;      TOPOLOGY:  linear
;    MOLECULE TYPE:  DNA (genomic)
US-08-757-669A-20

  Query Match             1.3%;  Score 54.6;  DB 3;  Length 32207;
  Best Local Similarity   56.4%;  Pred. No. 0.0022;
  Matches  102;  Conservative    0;  Mismatches   79;  Indels    0;  Gaps    0;

Qy          72 CTGCTCTTCCTGCTGTGGGTGCGCCTGAAGGGGCTGGAGTTCGTGCTCATCCACCAGCGC 131
               ||||||||||||||| |      | |||  |     |||   ||| ||||| ||||||  | |
Db       19294 CTGCTCTTCCTGCTCTTCCACCTCCTCTAACTCCTGCTCTTCCTGCTCTTCCACCTCCTC 19353

Qy         132 TGGGTGTTCGTGTGCCTCTTCCTCCTGCCGCTCTCGCTTATCTTCGATATCTACTACTAC 191
               |   | | |   |||  |||||||||  ||      |  || || |  ||  ||| | |
Db       19354 TAACTCCTGCTCCTCCTGCTCCTCCTGCTCCTGCTCTTGCTCCTCCACCTCCTCTAATTC 19413
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;      CLASSIFICATION:  435
;    ATTORNEY/AGENT INFORMATION:
;      NAME:  White, John P.
;      REGISTRATION NUMBER:  28,678
;      REFERENCE/DOCKET NUMBER:  0575/52268/JPW/MSC/SKS
;    TELECOMMUNICATION INFORMATION:
;      TELEPHONE:  212-278-0400
;      TELEFAX:  212-391-0525
;  INFORMATION FOR SEQ ID NO:  1:
;    SEQUENCE CHARACTERISTICS:
;      LENGTH:  3489 base pairs
;      TYPE:  nucleic acid
;      STRANDEDNESS:  single
;      TOPOLOGY:  linear
;    MOLECULE TYPE:  DNA (genomic)
;    FEATURE:
;      NAME/KEY:  CDS
;      LOCATION:  1..3489
US-08-728-323A-1

  Query Match             1.3%;  Score 54.6;  DB 2;  Length 3489;
  Best Local Similarity   56.4%;  Pred. No. 0.00058;
  Matches  102;  Conservative    0;  Mismatches   79;  Indels    0;  Gaps    0;

Qy         72 CTGCTCTTCCTGCTGTGGGTGCGCCTGAAGGGGCTGGAGTTCGTGCTCATCCACCAGCGC 131
              ||||||||||||||| |     | |||  |    |||   ||| ||||| ||||||  | |
Db       2703 CTGCTCTTCCTGCTCTTCCACCTCCTCTAACTCCTGCTCTTCCTGCTCTTCCACCTCCTC 2644

Qy        132 TGGGTGTTCGTGTGCCTCTTCCTCCTGCCGCTCTCGCTTATCTTCGATATCTACTACTAC 191
              |   |  |  |   |||  |||||||||  ||      |  || || |  ||  ||| | |
Db       2643 TAACTCCTGCTCCTCCTGCTCCTCCTGCTCCTGCTCTTGCTCCTCCACCTCCTCTAATTC 2584

Qy        192 GTGCGCGCCTGGGTGGTGTTCAAGCTCAGCAGCGCTCCGCGCCTGCACGAGCAGCGCGTG 251
               ||| |  |  | |  || ||  ||||  |  |   | || | ||| |   || | | ||
Db       2583 CTGCTCTTCCTGCTCCTGCTCTTGCTCTTCCACCTCCTGCTCTTGCTCTTCCACCTCCTG 2524

Qy        252 C 252
              |
Db       2523 C 2523


RESULT 9
US-09-298-568-1/c
; Sequence 1, Application US/09298568
; Patent No. 6322792
; GENERAL INFORMATION:
;  APPLICANT: Kieff, Elliott D.
;  APPLICANT: Ballestas, Mary E.
;  APPLICANT: Kaye, Kenneth M.
;  TITLE OF INVENTION: RHADINO VIRUS LANA ACTS IN TRANS ON A UNIT OF RHADINO
;  TITLE OF INVENTION: VIRUS DNA TO MEDIATE EFFICIENT EPISOME PERSISTENCE
;  FILE REFERENCE: 16412-10001R
;  CURRENT APPLICATION NUMBER: US/09/298,568
;  CURRENT FILING DATE: 1999-04-21
;  EARLIER APPLICATION NUMBER: US 60/109,422
;  EARLIER FILING DATE: 1998-11-19
;  NUMBER OF SEQ ID NOS: 3
;  SOFTWARE: PatentIn Ver. 2.0
; SEQ ID NO 1
;   LENGTH: 3489
;   TYPE: DNA
;   ORGANISM: Kaposi's sarcoma-associated herpesvirus
US-09-298-568-1

  Query Match             1.3%;  Score 54.6;  DB 3;  Length 3489;
  Best Local Similarity   56.4%;  Pred. No. 0.00058;
  Matches  102;  Conservative    0;  Mismatches   79;  Indels    0;  Gaps    0;

Qy         72 CTGCTCTTCCTGCTGTGGGTGCGCCTGAAGGGGCTGGAGTTCGTGCTCATCCACCAGCGC 131
              ||||||||||||||| |     | |||  |    |||   ||| ||||| ||||||  | |
Db       2703 CTGCTCTTCCTGCTCTTCCACCTCCTCTAACTCCTGCTCTTCCTGCTCTTCCACCTCCTC 2644

Qy        132 TGGGTGTTCGTGTGCCTCTTCCTCCTGCCGCTCTCGCTTATCTTCGATATCTACTACTAC 191
              |   |  |  |   |||  |||||||||  ||      |  || || |  ||  ||| | |
Db       2643 TAACTCCTGCTCCTCCTGCTCCTCCTGCTCCTGCTCTTGCTCCTCCACCTCCTCTAATTC 2584

Qy        192 GTGCGCGCCTGGGTGGTGTTCAAGCTCAGCAGCGCTCCGCGCCTGCACGAGCAGCGCGTG 251
               ||| |  |  | |  || ||  ||||  |  |   | || | ||| |   || | | ||
Db       2583 CTGCTCTTCCTGCTCCTGCTCTTGCTCTTCCACCTCCTGCTCTTGCTCTTCCACCTCCTG 2524

Qy        252 C 252
              |
Db       2523 C 2523


RESULT 10
US-09-410-399-1/c
; Sequence 1, Application US/09410399
; Patent No. 6482587
; GENERAL INFORMATION:
;  APPLICANT:  Robertson, Erle S.
;  APPLICANT:  Cotter, Murray A.
;  TITLE OF INVENTION:  Methods to Inhibit or Enhance the Binding of Viral DNA
;  TITLE OF INVENTION:  to Genomic Host DNA
;  FILE REFERENCE:  UM-03778
;  CURRENT APPLICATION NUMBER: US/09/410,399
;  CURRENT FILING DATE:  1999-10-01
;  NUMBER OF SEQ ID NOS:  6
;  SOFTWARE:  PatentIn Ver. 2.0
; SEQ ID NO 1
;   LENGTH: 3489
;   TYPE: DNA
;   ORGANISM: Kaposi's sarcoma-associated herpesvirus
US-09-410-399-1

  Query Match             1.3%;  Score 54.6;  DB 4;  Length 3489;
  Best Local Similarity   56.4%;  Pred. No. 0.00058;
  Matches  102;  Conservative    0;  Mismatches   79;  Indels    0;  Gaps    0;

Qy         72 CTGCTCTTCCTGCTGTGGGTGCGCCTGAAGGGGCTGGAGTTCGTGCTCATCCACCAGCGC 131
              ||||||||||||||| |     | |||  |    |||   ||| ||||| ||||||  | |
Db       2703 CTGCTCTTCCTGCTCTTCCACCTCCTCTAACTCCTGCTCTTCCTGCTCTTCCACCTCCTC 2644

Qy        132 TGGGTGTTCGTGTGCCTCTTCCTCCTGCCGCTCTCGCTTATCTTCGATATCTACTACTAC 191
              |   |  |  |   |||  |||||||||  ||      |  || || |  ||  ||| | |
Db       2643 TAACTCCTGCTCCTCCTGCTCCTCCTGCTCCTGCTCTTGCTCCTCCACCTCCTCTAATTC 2584

Qy        192 GTGCGCGCCTGGGTGGTGTTCAAGCTCAGCAGCGCTCCGCGCCTGCACGAGCAGCGCGTG 251
               ||| |  |  | |  || ||  ||||  |  |   | || | ||| |   || | | ||
Db       2583 CTGCTCTTCCTGCTCCTGCTCTTGCTCTTCCACCTCCTGCTCTTGCTCTTCCACCTCCTG 2524

Qy        252 C 252
              |
Db       2523 C 2523


RESULT 11
US-09-894-273-1/c
; Sequence 1, Application US/09894273
; Patent No. 6756203
; GENERAL INFORMATION:
;  APPLICANT: Kieff, Elliott D.
;  APPLICANT:  Ballestas, Mary E.
;  APPLICANT:  Kaye, Kenneth M.
;  TITLE OF INVENTION: RHADINO VIRUS LANA ACTS IN TRANS ON A UNIT OF RHADINO
;  TITLE OF INVENTION:  VIRUS DNA TO MEDIATE EFFICIENT EPISOME PERSISTENCE
;  FILE REFERENCE: 16412-10001R
;  CURRENT APPLICATION NUMBER: US/09/894,273
;  CURRENT FILING DATE:  2001-06-28
;  PRIOR APPLICATION NUMBER: US 60/109,422
;  PRIOR FILING DATE: 1998-11-19
;  NUMBER OF SEQ ID NOS: 3
;  SOFTWARE: PatentIn Ver. 2.0
; SEQ ID NO 1
;   LENGTH: 3489
;   TYPE: DNA
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;  PRIOR APPLICATION NUMBER: 60/237,768
;  PRIOR FILING DATE: 2000-10-03
;  PRIOR APPLICATION NUMBER: 60/231,498
;  PRIOR FILING DATE: 2000-09-08
;  NUMBER OF SEQ ID NOS: 207012
;  SOFTWARE: FastSEQ for Windows Version 4.0
; SEQ ID NO 16295
;   LENGTH: 140982
;   TYPE: DNA
;   ORGANISM: Human
US-09-949-016-16295

  Query Match             1.3%;  Score 55.8;  DB 4;  Length 140982;
  Best Local Similarity   61.2%;  Pred. No. 0.0025;
  Matches   90;  Conservative    0;  Mismatches   57;  Indels    0;  Gaps    0;

Qy       1766 AAGAATGTGGGATTTGGAGTCAGACAAGCCTGAGTCCAGTTCCCCGTTTAGAACTCATTA 1825
             || ||  | |||| | |||||||     |  || || ||||    || |   |  |||||
Db      81151 AAAAACATAGGATGTAGAGTCAGGAGCCCTGGATTCTAGTTTTGGGTCTTCCAGGCATTA 81092

Qy       1826 GCTGTGTGACTCTGGGTGAGTCCCTTAACCCCTCTGAGCCCGGGTCTCTTCATTAGTTGA 1885
             |||| || |   | ||  |||| ||||| |  | ||||  |  |||||||||| ||  |
Db      81091 GCTGAGTTATGTTAGGGAAGTCTCTTAAGCTATTTGAGATCCAGTCTCTTCATCCGTAAA 81032

Qy       1886 AAGGGATAGTAATACCTACTTGCAGGT 1912
              |   | | |||||    ||| |||||
Db      81031 TATACAAAATAATAAGGCCTTCCAGGT 81005


RESULT 6
US-09-949-016-162721/c
; Sequence 162721, Application US/09949016
; Patent No. 6812339
; GENERAL INFORMATION:
;  APPLICANT: VENTER, J. Craig et al.
;  TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES ASSOCIATED
;  TITLE OF INVENTION:  WITH HUMAN DISEASE, METHODS OF DETECTION AND USES THEREOF
;  FILE REFERENCE: CL001307
;  CURRENT APPLICATION NUMBER: US/09/949,016
;  CURRENT FILING DATE:  2000-04-14
;  PRIOR APPLICATION NUMBER: 60/241,755
;  PRIOR FILING DATE: 2000-10-20
;  PRIOR APPLICATION NUMBER: 60/237,768
;  PRIOR FILING DATE: 2000-10-03
;  PRIOR APPLICATION NUMBER: 60/231,498
;  PRIOR FILING DATE: 2000-09-08
;  NUMBER OF SEQ ID NOS: 207012
;  SOFTWARE: FastSEQ for Windows Version 4.0
; SEQ ID NO 162721
;   LENGTH: 601
;   TYPE: DNA
;   ORGANISM: Human
US-09-949-016-162721

  Query Match             1.3%;  Score 55.4;  DB 4;  Length 601;
  Best Local Similarity   60.5%;  Pred. No. 0.00012;
  Matches   89;  Conservative    1;  Mismatches   57;  Indels    0;  Gaps    0;

Qy       1766 AAGAATGTGGGATTTGGAGTCAGACAAGCCTGAGTCCAGTTCCCCGTTTAGAACTCATTA 1825
             || ||  | |||| | |||||||     |  || || ||||    || |   |  |||||
Db        416 AAAAACATAGGATGTAGAGTCAGGAGCCCTGGATTCTAGTTTTGGGTCTTCCAGGCATTA 357

Qy       1826 GCTGTGTGACTCTGGGTGAGTCCCTTAACCCCTCTGAGCCCGGGTCTCTTCATTAGTTGA 1885
             |||| || |   | ||  |||| ||||| |  | ||||  |  |||||||||| :| |
Db        356 GCTGAGTTATGTTAGGGAAGTCTCTTAAGCTATTTGAGATCCAGTCTCTTCATCCRTAAA 297

Qy       1886 AAGGGATAGTAATACCTACTTGCAGGT 1912
              |   | | |||||    ||| |||||
Db        296 TATACAAAATAATAAGGCCTTCCAGGT 270


RESULT 7
US-09-949-016-17215/c
; Sequence 17215, Application US/09949016
; Patent No. 6812339
; GENERAL INFORMATION:
;  APPLICANT: VENTER, J. Craig et al.
;  TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES ASSOCIATED
;  TITLE OF INVENTION:  WITH HUMAN DISEASE, METHODS OF DETECTION AND USES THEREOF
;  FILE REFERENCE: CL001307
;  CURRENT APPLICATION NUMBER: US/09/949,016
;  CURRENT FILING DATE:  2000-04-14
;  PRIOR APPLICATION NUMBER: 60/241,755
;  PRIOR FILING DATE: 2000-10-20
;  PRIOR APPLICATION NUMBER: 60/237,768
;  PRIOR FILING DATE: 2000-10-03
;  PRIOR APPLICATION NUMBER: 60/231,498
;  PRIOR FILING DATE: 2000-09-08
;  NUMBER OF SEQ ID NOS: 207012
;  SOFTWARE: FastSEQ for Windows Version 4.0
; SEQ ID NO 17215
;   LENGTH: 23193
;   TYPE: DNA
;   ORGANISM: Human
;   FEATURE:
;   NAME/KEY: misc_feature
;   LOCATION: (1)...(23193)
;   OTHER INFORMATION: n = A,T,C or G
US-09-949-016-17215

  Query Match             1.3%;  Score 55.2;  DB 4;  Length 23193;
  Best Local Similarity   63.6%;  Pred. No. 0.0012;
  Matches   84;  Conservative    0;  Mismatches   48;  Indels    0;  Gaps    0;

Qy       1759 GAATGGAAAGAATGTGGGATTTGGAGTCAGACAAGCCTGAGTCCAGTTCCCCGTTTAGAA 1818
             ||| ||  || ||   || |  |||     ||||  |||||||     |||| | |  |
Db      14460 GAAGGGGCAGCATAGAGGCTCAGGAACTGCACAAAGCTGAGTCTGCATCCCAGCTCTGCC 14401

Qy       1819 CTCATTAGCTGTGTGACTCTGGGTGAGTCCCTTAACCCCTCTGAGCCCGGGTCTCTTCAT 1878
              | ||||||||| |||||  || |  |||| ||| ||| |||||||| |    | || ||||
Db      14400 TTTATTAGCTGGGTGACCTTGAGCAAGTCACTTCACCTCTCTGAGCTCCAATTTCCTCAT 14341

Qy       1879 TAGTTGAAAGGG 1890
              | ||  || |||
Db      14340 TGGTAAAATGGG 14329


RESULT 8
US-08-728-323A-1/c
; Sequence 1, Application US/08728323A
; Patent No. 5948676
;  GENERAL INFORMATION:
;    APPLICANT:  Chang, Yuan
;    APPLICANT:  Bohenzky, Roy A.
;    APPLICANT:  Russo, James J.
;    APPLICANT:  Edelman, Isidore S.
;    APPLICANT:  Moore, Patrick S.
;    TITLE OF INVENTION:  Immediate Early Protein From Kaposi's
;    TITLE OF INVENTION:  Sarcoma-Associated Herpesvirus, DNA
;    TITLE OF INVENTION:  Encoding Same And Uses Thereof
;    NUMBER OF SEQUENCES:  21
;    CORRESPONDENCE ADDRESS:
;      ADDRESSEE:  Cooper & Dunham LLP
;      STREET:  1185 Avenue of the Americas
;      CITY:  New York
;      STATE:  New York
;      COUNTRY:  U.S.A.
;      ZIP:  10036
;    COMPUTER READABLE FORM:
;      MEDIUM TYPE:  Floppy disk
;      COMPUTER:  IBM PC compatible
;      OPERATING SYSTEM:  PC-DOS/MS-DOS
;      SOFTWARE:  PatentIn Release #1.0, Version #1.30
;    CURRENT APPLICATION DATA:
;      APPLICATION NUMBER:  US/08/728,323A
;      FILING DATE:
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; Sequence 2284, Application US/09513999C
; Patent No. 6783961
; GENERAL INFORMATION:
;  APPLICANT: Dumas Milne Edwards, J.B.
;  APPLICANT:  Duclert, A.
;  APPLICANT:  Giordano, J.Y.
;  TITLE OF INVENTION: Expressed Sequence Tags and Encoded Human Proteins.
; Patent No. 6783961
;  FILE REFERENCE: 59.US2.REG
;  CURRENT APPLICATION NUMBER: US/09/513,999C
;  CURRENT FILING DATE:  2000-02-24
;  PRIOR APPLICATION NUMBER: US 60/122,487
;  PRIOR FILING DATE: 1999-02-26
;  NUMBER OF SEQ ID NOS: 36681
;  SOFTWARE: Patent.pm
; SEQ ID NO 2284
;   LENGTH: 424
;   TYPE: DNA
;   ORGANISM: Homo sapiens
;   FEATURE:
;   NAME/KEY: CDS
;   LOCATION: 178..423
;   FEATURE:
;   NAME/KEY: misc_feature
;   LOCATION: 126
;   OTHER INFORMATION: k=g or t
;   FEATURE:
;   NAME/KEY: misc_feature
;   LOCATION: 381
;   OTHER INFORMATION: n=a, g, c or t
US-09-513-999C-2284

  Query Match             3.7%;  Score 156;  DB 4;  Length 424;
  Best Local Similarity   98.8%;  Pred. No. 2.6e-32;
  Matches  167;  Conservative    0;  Mismatches    1;  Indels    1;  Gaps    1;

Qy       4018 CAGTGTATGGTGTGTGATTTTTATTTTCTTCTTTGGTCATGGGGGCCAAGGAGAAAGGCA 4077
             ||||||||||||||||||||||||||||||||||||||||| ||| ||||||||||||||||
Db        424 CAGTGTATGGTGTGTGATTTTTATTTTCTTCTTTGGTCAT-GGGNCCAAGGAGAAAGGCA 366

Qy       4078 TGAATCTTCCCTGTCAGGCTCTTACAGCCACAGGCACTGTGTCTACTGTCTGGAAGACAT 4137
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        365 TGAATCTTCCCTGTCAGGCTCTTACAGCCACAGGCACTGTGTCTACTGTCTGGAAGACAT 306

Qy       4138 GTCCCCGTGGCTGTGGGGCCGCTGCTTCTGTTTAAATAAAAGTGGCCTG 4186
             ||||||||||||||||||||||||||||||||||||||||||||||||||
Db        305 GTCCCCGTGGCTGTGGGGCCGCTGCTTCTGTTTAAATAAAAGTGGCCTG 257


RESULT 3
US-09-949-016-162720/c
; Sequence 162720, Application US/09949016
; Patent No. 6812339
; GENERAL INFORMATION:
;  APPLICANT: VENTER, J. Craig et al.
;  TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES ASSOCIATED
;  TITLE OF INVENTION:  WITH HUMAN DISEASE, METHODS OF DETECTION AND USES THEREOF
;  FILE REFERENCE: CL001307
;  CURRENT APPLICATION NUMBER: US/09/949,016
;  CURRENT FILING DATE:  2000-04-14
;  PRIOR APPLICATION NUMBER: 60/241,755
;  PRIOR FILING DATE: 2000-10-20
;  PRIOR APPLICATION NUMBER: 60/237,768
;  PRIOR FILING DATE: 2000-10-03
;  PRIOR APPLICATION NUMBER: 60/231,498
;  PRIOR FILING DATE: 2000-09-08
;  NUMBER OF SEQ ID NOS: 207012
;  SOFTWARE: FastSEQ for Windows Version 4.0
; SEQ ID NO 162720
;   LENGTH: 601
;   TYPE: DNA
;   ORGANISM: Human
US-09-949-016-162720

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  Best Local Similarity   61.2%;  Pred. No. 9.5e-05;
  Matches   90;  Conservative    0;  Mismatches   57;  Indels    0;  Gaps    0;

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Qy       1826 GCTGTGTGACTCTGGGTGAGTCCCTTAACCCCTCTGAGCCCGGGTCTCTTCATTAGTTGA 1885
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Db        443 GCTGAGTTATGTTAGGGAAGTCTCTTAAGCTATTTGAGATCCAGTCTCTTCATCCGTAAA 384

Qy       1886 AAGGGATAGTAATACCTACTTGCAGGT 1912
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Db        383 TATACAAAATAATAAGGCCTTCCAGGT 357


RESULT 4
US-09-949-016-11777/c
; Sequence 11777, Application US/09949016
; Patent No. 6812339
; GENERAL INFORMATION:
;  APPLICANT: VENTER, J. Craig et al.
;  TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES ASSOCIATED
;  TITLE OF INVENTION:  WITH HUMAN DISEASE, METHODS OF DETECTION AND USES THEREOF
;  FILE REFERENCE: CL001307
;  CURRENT APPLICATION NUMBER: US/09/949,016
;  CURRENT FILING DATE:  2000-04-14
;  PRIOR APPLICATION NUMBER: 60/241,755
;  PRIOR FILING DATE: 2000-10-20
;  PRIOR APPLICATION NUMBER: 60/237,768
;  PRIOR FILING DATE: 2000-10-03
;  PRIOR APPLICATION NUMBER: 60/231,498
;  PRIOR FILING DATE: 2000-09-08
;  NUMBER OF SEQ ID NOS: 207012
;  SOFTWARE: FastSEQ for Windows Version 4.0
; SEQ ID NO 11777
;   LENGTH: 140925
;   TYPE: DNA
;   ORGANISM: Human
US-09-949-016-11777

  Query Match             1.3%;  Score 55.8;  DB 4;  Length 140925;
  Best Local Similarity   61.2%;  Pred. No. 0.0025;
  Matches   90;  Conservative    0;  Mismatches   57;  Indels    0;  Gaps    0;

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             || ||  | |||| | |||||||      |  || || ||||    || |   | |||||
Db      81151 AAAAACATAGGATGTAGAGTCAGGAGCCCTGGATTCTAGTTTTGGGTCTTCCAGGCATTA 81092

Qy       1826 GCTGTGTGACTCTGGGTGAGTCCCTTAACCCCTCTGAGCCCGGGTCTCTTCATTAGTTGA 1885
             |||| || |   | ||  |||| ||||| |  | ||||  |  |||||||||| ||  |
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Qy       1886 AAGGGATAGTAATACCTACTTGCAGGT 1912
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Db      81031 TATACAAAATAATAAGGCCTTCCAGGT 81005


RESULT 5
US-09-949-016-16295/c
; Sequence 16295, Application US/09949016
; Patent No. 6812339
; GENERAL INFORMATION:
;  APPLICANT: VENTER, J. Craig et al.
;  TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES ASSOCIATED
;  TITLE OF INVENTION:  WITH HUMAN DISEASE, METHODS OF DETECTION AND USES THEREOF
;  FILE REFERENCE: CL001307
;  CURRENT APPLICATION NUMBER: US/09/949,016
;  CURRENT FILING DATE:  2000-04-14
;  PRIOR APPLICATION NUMBER: 60/241,755
;  PRIOR FILING DATE: 2000-10-20
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GenCore version 5.1.6

OM nucleic - nucleic search, using sw model

Run on: August 17, 2005, 13:10:42 ; Search time 631.158 Seconds
(without alignments)
10854.792 Million cell updates/sec

Title: US-09-996-630A-10
Perfect score: 4187
Sequence: 1 ggcgcgaacccgcagcgctt.....tttaaataaaagtggcctgg 4187

Scoring table: IDENTITY_NUC
Gapop 10.0 , Gapext 1.0

Searched: 1202784 seqs, 818138359 residues

Total number of hits satisfying chosen parameters: 2405568

Minimum DB seq length: 0
Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries

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6: /cgn2_6/ptodata/1/ina/backfiles1.seq:*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	% Query Match	Length	DB	ID	Description
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c 3	55.8	1.3	601	4	US-09-949-016-162720	Sequence 162720,
c 4	55.8	1.3	140925	4	US-09-949-016-11777	Sequence 11777, A
c 5	55.8	1.3	140982	4	US-09-949-016-16295	Sequence 16295, A
c 6	55.4	1.3	601	4	US-09-949-016-162721	Sequence 162721,
c 7	55.2	1.3	23193	4	US-09-949-016-17215	Sequence 17215, A
c 8	54.6	1.3	3489	2	US-08-728-323A-1	Sequence 1, Appli
c 9	54.6	1.3	3489	3	US-09-298-568-1	Sequence 1, Appli
c 10	54.6	1.3	3489	4	US-09-410-399-1	Sequence 1, Appli
c 11	54.6	1.3	3489	4	US-09-894-273-1	Sequence 1, Appli
12	54.6	1.3	32207	2	US-08-770-379-20	Sequence 20, Appl
13	54.6	1.3	32207	3	US-08-757-669A-20	Sequence 20, Appl
14	54.6	1.3	32207	3	US-09-230-371A-20	Sequence 20, Appl
15	54.2	1.3	262	4	US-09-573-080A-99	Sequence 99, Appl
16	54.2	1.3	92681	4	US-09-949-016-14772	Sequence 14772, A
17	53.8	1.3	29321	4	US-09-949-016-14257	Sequence 14257, A
18	53.8	1.3	29321	4	US-09-949-016-14258	Sequence 14258, A
19	53.8	1.3	55298	4	US-09-491-356C-1	Sequence 1, Appli
c 20	53.6	1.3	66219	4	US-09-949-016-12038	Sequence 12038, A
c 21	53.6	1.3	66227	4	US-09-949-016-15303	Sequence 15303, A
c 22	52.8	1.3	256287	4	US-09-949-016-14608	Sequence 14608, A
23	52	1.2	601	4	US-09-949-016-67372	Sequence 67372, A
24	52	1.2	601	4	US-09-949-016-67442	Sequence 67442, A
25	52	1.2	601	4	US-09-949-016-67512	Sequence 67512, A
c 26	52	1.2	49931	4	US-09-949-016-13727	Sequence 13727, A
c 27	52	1.2	49931	4	US-09-949-016-13728	Sequence 13728, A
c 28	52	1.2	49931	4	US-09-949-016-13729	Sequence 13729, A
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33	50.8	1.2	92155	4	US-09-949-016-17484	Sequence 17484, A
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37	50.4	1.2	30000	4	US-10-007-010-10	Sequence 10, Appl
38	50.4	1.2	53562	4	US-09-949-016-16286	Sequence 16286, A
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c 40	50.4	1.2	144596	4	US-09-949-016-13035	Sequence 13035, A
c 41	50.2	1.2	64309	4	US-09-949-016-14581	Sequence 14581, A
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c 44	50	1.2	110266	4	US-09-949-016-14914	Sequence 14914, A
c 45	50	1.2	110266	4	US-09-949-016-14915	Sequence 14915, A

ALIGNMENTS

RESULT 1
US-09-313-294A-6606
; Sequence 6606, Application US/09313294A
; Patent No. 6476212
; GENERAL INFORMATION:
; APPLICANT: Lalgudi, Raghunath V.
; APPLICANT: Ito, Laura Y.
; APPLICANT: Sherman, Bradley K.
; TITLE OF INVENTION: POLYNUCLEOTIDES AND POLYPEPTIDES DERIVED FROM CORN EAR
; FILE REFERENCE: PL-0017 US
; CURRENT APPLICATION NUMBER: US/09/313,294A
; CURRENT FILING DATE: 1999-05-14
; NUMBER OF SEQ ID NOS: 7600
; SOFTWARE: PERL Program
; SEQ ID NO 6606
; LENGTH: 275
; TYPE: DNA
; ORGANISM: Zea mays
; FEATURE:
; NAME/KEY: misc_feature
; OTHER INFORMATION: Incyte ID No. 6476212 700352091H1
US-09-313-294A-6606

Query Match 4.9%; Score 206.4; DB 4; Length 275;
Best Local Similarity 86.6%; Pred. No. 3.1e-46;
Matches 239; Conservative 0; Mismatches 36; Indels 1; Gaps 1;

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Qy       1269 TGGCTGTGTCCGTTCATCCTGCCCAGCCAGCCAGGCCTAGTGCACCCCAAAGGAAATGAG 1328
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Qy       1449 TATGCCGACTGCTACATGAACCGGGAGGAGTTCTGG 1484
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RESULT 2
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